

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number

TO: Minh-Tam Davis

Location: REM-3A24&3C18

Art Unit: 1642

Thursday, March 09, 2006

Case Serial Number: 10/001469

From: Alex Waclawiw

Location: Biotech-Chem Library

Rem 1A71

Phone: 272-2534

Alexandra.waclawiw@uspto.gov

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STIC-Biotech/ChemLib

181717

From:

Chan, Christina

Sent: To: Wednesday, March 08, 2006 2:25 PM Davis, Minh-Tam; STIC-Biotech/ChemLib

Subject:

RE: Rush search request for 10/001469

Please rush. Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 (571)-272-0841 Remsen, 3E89 M

----Original Message----

From:

Davis, Minh-Tam

Sent:

Wednesday, March 08, 2006 11:19 AM

To:

Chan, Christina

Subject:

Rush search request for 10/001469

Please search in commercial database, issued patent files, PGPUB and interference:

The protein of SEQ ID NO:2866

Thank you.

MINH TAM DAVIS

ART UNIT 1642, ROOM 3A24, MB 3C18

272-0830

Point of Contact:

Alexandra Waclawiw
Searcher:

Searcher Phone:
Date Searcher Picked up:
Date completed:
Searcher Prep Time:
Online Time:

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Type of	Search			
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The invention relates to a novel composition comprising: a substance that modulates the status of a protein comprising a sequence with a fully defined 2466 or 3136 amino acid sequence given in the specification; or a molecule that is immobilised by a protein comprising a sequence with a fully defined 2466 or 3136 amino acids, where the status of a cell expressing the protein is modulated. The invention further relates to: an antibody; a non-human transgenic animal or hybridoma that produces the antibody; a method of delivering a cytotoxic or diagnostic agent to a cell that expresses the protein; a polynucleotide that encodes the protein; a method for inhibiting the growth of cancer cells that express the protein; a method for generating a mammalian immune response directed to the protein; a method detecting in a sample the presence of a 10193All
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14-DEC-2001; 2001US-00017666
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Matches 318;
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New composition comprising 101P3A11-related protein, useful preventing or treating cancer e.g., stomach, cervix, uterus, prostate, colon, kidney or breast cancer.
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31-OCT-2001; 2001US-00001469.
14-DEC-2001; 2001US-00017666.
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31-OCT-2001;
14-DEC-2001;
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06-OCT-1998;
06-OCT-1998;
11-MAY-1999;
                                                                       Tang YT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
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                                                                                                                                                                  13-APR-2000.
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                                                                                                 (INCY-)
                                                                 Y YT,
                                                2000-328934/28.
DB; AAA09351.
                                                                                                 INCYTE PHARM INC.
                                                                      Yue H, Lal P,
Guegler KJ,
                                                                                                                98US-00167219.
98US-0172211P.
99US-0133585P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunostimulant.
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                                                                                                                                                 99WG-US023317.
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57
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272
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    "G-protein_coupled :
                                                                                                                                                                                                 "potential phosphorylation site"
                                                                                                                                                                                                                  "potential phosphorylation site"
                                                                                                                                                                                                                                 "G-protein coupled receptor"
                                                                                                                                                                                                                                                                                                  "G-protein coupled receptor"
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                                                                        Bandman O,
Gorgone GA,
                                                                                                                                                           0002/62/11
                                                                                                                                                                                                                                                                                                                                           glycosylation site"
                                                                        Au-Young J
Baughn MR,
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د.
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Novel human G-protein coupled receptor proteins used in the diagnosis, treatment and prevention of nervous system disorders, autoimmune/inflammatory disorders, and cell proliferative disorders su

disorders such

as cancer.

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Best Local S
Matches 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence encodes human G-protein coupled receptor protein (GCRP) The GCRP polypeptides, polymucleotides, antibodies, antagonists and agonists may be administered to human patients for the diagnosis, treatment and prevention of nervous system disorders (e.g. epilepsy, stroke, neoplasms, Alzheimer's disease), autoimmune or inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disorders, complications of cancer, hemadialysis and extracorporeal circulation, and cell proliferative disorders. They are also used to treat or prevent disorders associated with decreased or increased
                   Xu J, Dillon DC, Mitcham Kalos MD, Fanger GR, Day Wang A, Meagher MJ;
WPI; 2001-425873/45
                                                                                            14-JAN-2000; 2000US-00483672.
                                                                                                                   16-JAN-2001; 2001WO-US001574.
                                                                                                                                                                                                              Human; prostate cancer; prostate-specific; diagnosis; vaccine; cytostatic; gene therapy; metastasis.
                                                                                                                                                                                                                                                  P835P amino acid sequence
                                                                                                                                                                                                                                                                          04-OCT-2001
                                                                                                                                                                                                                                                                                                                          AAM01306 standard;
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                                                                     (CORI-) CORIXA CORP.
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317; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                               GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKE 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MAPDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS
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                                                                                                                                                                                                                                                                          (first entry)
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99.7%;
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Pred. No. 2.2e-169;
1; Mismatches 0;
                                    Harlocker SL, Jiang
Retter MW, Stolk JA,
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Claim 2; Page 534-535; 543pp; English
                                                                                                                            New polynucleotide encoding a prostate-specific protein, monitoring and treating prostate cancer in a patient and
                                                                                                                             for diagnosing, for use in
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The present invention describes polynucleotide sequences (I) which encode CC prostate-specific proteins (II). (I) and (II) have cytostatic activity, CC and can be used in vaccine production and gene therapy. (I), (II), CC antibodies to (II), fusion proteins comprising (II), and isolated T cells CC prepared using (I) or (II) are used treat cancer in a patient. (I) and CC cancer that is diagnosed or treated is particularly prostate cancer. The CC and (II) can be used in vaccines. The antibodies or (I) can be used for CC concer that progression of cancer in a patient. (I) and (II) can be used for concer in a patient. (I) and (II) can also CC be used to improve diagnostic and therapeutic methods for prostate CC volume. AAH93357 to AAH93944 and AAM01115 to AAM01318 represent CC colynucleotide and amino acid sequences used in the exemplification of the present invention

Sequence 318 AA,

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317; Conserv
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              IRQRILRLFHVATHASEP 318
                                                              GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKE 300
                                                                                                               YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF
                                                                                                                                                                 MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS
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                                                 GTCVSHVCAVF1FYVPF1GLSMVHRFSKRRDSPLPV1LAN1YLLVPPVLNP1VYGVKTKB
                                                                                                                                                                                                                                                                                                   99.8%; Score 1619; DB 4; ilarity 99.7%; Pred. No. 2.2e-169; Conservative 1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                           Length 318;
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AAU69951 standard; protein; 318

AAU69951;

30-JAN-2002 (first entry)

Human prostate cDNA encoded protein

Human; prostate cancer; cytostatic; immunostimulant; tumour; immunogen.

WO200173032-A2

AAUG9951 ID AAUG9951 AC AAUG AC AAUG XX AAUG DT 30-J DT 30-J XX Huma XX Huma XX Huma XX Homc OS Homc YX WO20 YX WO20 YX Y 04-OCT-2001.

27-MAR-2001; 2001WO-US009919

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RESULT 8
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09-MAY-2000
12-MAY-2000
13-JUN-2000
27-JUN-2000
09-AUG-2000
09-AUG-2000
06-SEP-2000
02-CCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to isolated prostate-specific polynucleotides, polypeptides, fusion proteins of the polypeptides, antibodies raised against the polypeptides (or antigenic epitopes derived from them) and antigen-presenting cells expressing the polypeptides. The antibodies are useful for detecting the presence of cancer, especially prostate cancer. The polypeptides, polynucleotides and the antigen-presenting cells are useful for stimulating and/or expanding T cells specific for a tumour protein, and for inhibiting the development of cancer especially prostate cancer. Compositions comprising the polynucleotide and/or polypeptide are useful for stimulating an immune response, and for treating cancer. The oligonucleotide is useful for detecting cancer. The present sequence is a prostate specific polypeptide of the invention
 AAG71659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 318 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-639232/73.
N-PSDB; AAS64174, AAS64176, AAS64177.
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Li SX,
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GR, Retter MW, Stolk JA,
Wang A, Skeiky YAW, Hep.
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standard; protein; 318
                                                                       IRQRILRLFHVATHASEP
                                                                                                                                                   GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSFLPVILANIYLLVPFVLNPIVYGVKTKE 300
                                                                                                                                                                                                          YCLHQDVWKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF 240
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2000US-00568100.
2000US-00570737.
2000US-00593793.
2000US-006362783.
2000US-00636215.
2000US-00657279.
2000US-00679426.
2000US-00685166.
2000US-00685166.
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99.7%;
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                                                                                                   318
                                                                                                                                                                                                                                                                                                                                                                                                                              1;
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Pred. No. 2.2e-169;
1; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 318;
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Carter D;
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Matches 317;
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                                                                                                                                                                                                                                                                                                                Sequence 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 11; Page 826-827; 1857pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polynucleotides which sensation for identifying
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24-FEB-2000; 2000US-0184809P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human olfactory receptor polypeptide,
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                                                                                                                                                                       PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA
             GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLFVILANIYLLVPPVLNPIVYGVKTKE
                                                                                                     MAFDRYVAI CHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS
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Pred. No. 2.2e-169;
1; Mismatches 0;
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and antagonists.
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RESULT 9
AABO2495
ID AABO
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XX AABO
XX AABO
XX AABO
XX Huma
XX Huma
XX Huma
XX Huma
XX Homx
XX H
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28-OCT-1999;
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28-OCT-1999;
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                                                                                                                                                                                                                                                     03-MAY-2001
                                                                                                                                                                                                                                                                                                  WO200131014-A2.
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99US-00427653.
99US-00427859.
99US-00428020.
99US-00428114.
99US-00429517.
99US-00429555.
99US-00429676.
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/note= "Third EC loop"
274. .295
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/label= Intracellular domain
/note= "First IC loop"
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/note= "Second EC loop"
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/note= "First EC loop"
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"Third IC loop"
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"Second IC loop"
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RESULT 10
ABB44533
ID ABB44
XX
AC ABB44
XX
DT 28-JA
DX
DE Human
XX
KW Human

ABB44533 standard;

protein;

ABB44533; 28-JAN-2002

(first entry)

Human; GCPR; G-coupled protein-receptor; cardiant; antiarteriosclerotic;

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Human GPCR7a polypeptide SEQ ID NO

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches 317;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to human G protein-coupled receptor (GPCR) and their corresponding DNA molecules. GPCR is also referred as seven transmembrane receptor. G protein-coupled receptor protein is useful for treating neurological disorder, particularly schizophrenia. GPCR protein is also useful for identifying compounds useful for treating schizophrenia. These compounds are also useful for treating other neurological and psychiatric diseases, e.g. depression, anxiety, bipolar disease, affective disorder, ettention deficit hyperactivity disorder/attention deficit disorder, epilepsy, neuritis, neurasthenia, neuropathy, neurosis, Alzheimer's disease, Parkinson's disease, migraine and senile dementia. The invention also provides genetic screening procedures that entail analysing a person's genome with respect to GPCR. The vectors are useful for the recombinant production of the GPCR's. The present sequence is human CON198 G protein-coupled receptor (GPCR) protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seven transmembrane receptor polypeptides and polynucleotides, useful iteating neurological or psychiatric disorders, e.g. schizophrenia, as well as for identifying compounds useful for treating schizophrenia.
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N-PSDB; AAD06504.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-DEC-1999; 99US-00454399.
12-JAN-2000; 2000US-00481794.
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                                                                   GTCVSHVCAVFIFYVPFIGLSMYHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKB
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IRQRILRLFHVATHASEP 318
                      IRQRILRLFHVATHASEP 318
                                                                                                                                          YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF
                                                                                                                                                                 YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGITREAQAKAF
                                                                                                                                                                                                                                                   MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS 180
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nilarity 99.7%;
Conservative
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Pred. No. 2.2e-169;
1; Mismatches 0;
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The invention relates to nucleic acid sequences (ABAS1529-ABAS1522) that encode G-coupled protein-receptor related polypeptides (ABB44522-CC ABB4453). The isolated polypeptide having a sequence differing by no CC more than 15 % of amino acid residues from one of 22 amino acid sequences (or mature forms of the sequences), fully defined in the specification CC and corresponding to human G-protein coupled receptor X (GPCEX) CC polypeptides. The polypeptides have potential cardiant, CC antiarteriosclerotic, anabolic, cytostatic and antiviral activity. The CC polypeptides can be administered theraputically, especially using gene CC therapy and expressing the encoding DNA in vivo, to treat or prevent CC GPCEX-associated disorders, especially in humans. For example, they can be used to treat/prevent cardiomyopathy, atherosclerosis, disorders (crelated to signal processing and metabolic pathway modulation (e.g. cobssity, anorexia), diabetes, osteoporosis, Crohn's disease, multiple collected to signal processing and metabolic pathway modulation (e.g. Alzheimer's disease, Patkinson's disease, neurodegenerative disorders, hamman edisorders, haematopoietic disorders, developmental diseases, neurological disorders, bacterial, fungal, proteozoal and viral infections (e.g. with human commondeficiency virus (HIV)-1 or HIV-2). They can be used disgnostically to determine the presence of or predisposition to a disease associated with altered levels of the polypeptide in mammals (especially humans) by detecting alterations in polypeptide in mammals (especially humans) by careful to identify agents binding polypeptide (e.g. cellular receptors or downstream effectors) and/or agents
         31-MAR-2000

05-APR-2000

06-APR-2000

06-APR-2000

06-APR-2000

06-APR-2000

06-APR-2000

06-APR-2000

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07-JUL-2000

07-JUL-2000

08-JUL-2000

09-APR-2000

09-APR-2000
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Padigaru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human G-protein coupled receptor X, GPCRX, polypeptide useful in treatment or prevention of GPCRX associated disorders e.g. cardiomyopathy or atherosclerosis, and to screen for antagonists and agonists useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 anabolic; cytostatic; antiviral; gene therapy; cardiomyopathy; obesity;
anorexia; diabetes; osteoporosis; Crohn's disease; multiple sclerosis;
asthma; Alzeimer's disease; Parkinson's disorder; Huntington's disease;
infection; human immunodeficiency virus; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       therapeutically.
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DB; ABA81542.
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2000US-0195068P.
2000US-0195069P.
2000US-0195070P.
2000US-0195170P.
2000US-0219859P.
2000US-0221325P.
2000US-0221325P.
2000US-0221325P.
2000US-0263638P.
2001US-02636333P.
2001US-02636433P.
2001US-0263644P.
2001US-0263644P.
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Mishnu VS,
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Tchernev VT,
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Spytek KA,
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Li L, Baumgartner
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RESULT 11
ABUT1842
ID 71842
ID 7847
AC ABUT1
XX ABUT1
XX Prost
XX PSMA.
XX PSMA.
XX PSMA.
XX PSMA.
XX WS20
PR 19-DI
XX 04-O
PR 29-UI
XX XX (XUJ)
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Best Local
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04-OCT-2000; 2000US-00679272
28-MAR-2001; 2001US-00822827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Prostate cancer; vaccine; gene therapy; cytostatic; fusion protein; immunogen; cancer; prostate specific antigen; PSA; prostatic acid phosphatase; PAP; prostate specific membrane antigen
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                                                                                                                                                                                           (XUJJ/)
(MITC/)
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                    MITCHAM J L.
HARLOCKER S.
JIANG Y.
KALOS M D.
FANGER G R.
RETTER M W.
STOLK J A.
DAY C H.
VEDVICK T S.
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Pred. No. 2.2e
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Li SX, Wang
Mcneill PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a fusion protein comprising at least one amino acid sequence of immunogenic portions of any of the 3 sequences not defined in the specification, or sequences having at least 70 or 90 % sequence identity to any one of the 35 sequences defined in the USPTO we site, which is encoded by any of the 4 nucleotide sequences not defined in the specification. The fusion protein, composition and methods are useful for diagnosing, preventing and/or treating cancer, particularly prostate cancer. The proteins are useful as markers to indicate the presence or absence of cancer. This is the amino acid sequence of a prostate cancer therapy associated protein. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from the US patent office at
  AAU24561;
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HOUGHTON R L.
Y DE BASSOLS C
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HEPLER W T.
HENDERSON R A.
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                            standard;
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Houghton RL,
                           protein; 318
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24-MAR-2000;
12-APR-2000;
12-APR-2000;
24-APR-2000;
26-MAY-2000;
23-JUN-2000;
                                                                                                                                                                                                                                                                                                                       The invention relates to nucleic acids encoding human olfactory receptors, OR, (a G protein-coupled receptor, GPCR). The OR's specifically recognise molecules, odourants, that elicit specific olfactory sensation. The human olfactory receptors and polynucleotides encoding them are useful for screening a library of chemical compounds for compounds that are involved in olfactory sensation. Modulators of their activity are useful for pharmacological and genetic modulation of olfactory signalling pathways. Therefore, they can be used in the food, pharmaceutical and cosmetic industries to customise odours and fragrances. The present sequence is a human olfactory receptor of the
                                                                                                                                                                                                                                                                                  Sequence 318
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07-SEP-2000;
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2000US-0192033P.

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17 2000US-019933SP.

17 2000US-0207702P.

17 2000US-0213849P.

17 2000US-0236734P.

17 2000US-0236732P.

17 2001US-0266862P.
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2.2e-169;
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01-AUG-1997; 97US-00904804.
10-FEB-1998; 98US-00020956.
25-FEB-1998; 98US-00115453.
23-SEP-1998; 98US-001232149.
15-JAN-1999; 99US-00232149.
15-JAN-1999; 99US-00232149.
16-AUG-1999; 99US-00232616.
13-UI-1999; 99US-00443686.
13-UI-1999; 99US-00443686.
13-UI-1999; 99US-00443686.
13-WAY-2000; 2000US-00536857.
29-WAY-2000; 2000US-005580100.
12-MAY-2000; 2000US-005580100.
12-MAY-2000; 2000US-005580100.
12-MAY-2000; 2000US-005580100.
12-MAY-2000; 2000US-005580100.
12-MAY-2000; 2000US-005580100.
12-MAY-2000; 2000US-00557037.
13-UUN-2000; 2000US-00557236.
06-SEP-2000; 2000US-00657236.
06-SEP-2000; 2000US-00657236.
06-SEP-2000; 2000US-00657279.
02-OCT-2000; 2000US-00657279.
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                                                    (WANG/)
(SKEI/)
                                  (HEND/)
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                                                CARTER D.

LI S X.

WANG A.

SKEIKY Y A W.

HEPLER W T.
                                                                                                                                 FANGER G R.
RETTER M W.
STOLK J A.
DAY C H.
VEDVICK T S.
 Dillon DC,
                                                                                                                                                                                                                   JIANG Y.
KALOS M D.
                                                                                                                                                                                                                                                                    MITCHAM J L
                                                                                                                                                                                                                                                                                                       XU J.
                                                                                                                                                                                                                                                    MITCHAM J L.
HARLOCKER S L.
                                 HENDERSON R A.
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Mitcham JL,
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THE REPORT OF THE PROPERTY OF 
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Best Local Similarity 99.7%;
Matches 317; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides prostate-specific coding sequences and their encoded proteins. These can be used in the diagnosis and treatment of cancers, particularly prostate cancer. The present sequence is a protein described in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; GPCR; G protein coupled receptor; signal transduction; olfactory; drug development; gustatory; taste; fragrance; receptor.
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Li SX, Wang A, Skeiky YAW, Hepler WT, H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 318 AA;
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                                                                   04-AUG-2000; 2000JP-00237818.
13-FEB-2001; 2001JP-00034434.
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(NISC-) JAPAN SCI & TECHNOLOGY CORP.
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Pred. No. 2.2e-169;

    Vedvick TS,
Henderson RA;

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Best Local
                                                Human; olfactory and pheromone G protein coupled; receptor; GPCR; tranquilizer; antidepressant; neuroleptic; endocrine; anabolic; anorectic; taste; fragrance; food additive; cosmetic; cell migration; sterility; psychotic disorder; neurological disorder; anxiety; schizophrenia; manic depression; depression; axonal growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABP95942) by extracting open-reading frames containing 6-8 transmembrane domains with 250-1000 amino acid residues to give a gene homologous with a known GPCR gene. The receptor proteins and encoded genes are useful for studying in vivo signal transduction mechanism and identifying targets for drug development e.g. based on olfactory and gustatory receptors in form of agonists and antagonists by screening intrinsic and extrinsic ligands as bitter taste inhibitors, taste enhancers and fragrance improvers. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                          Human olfactory and pheromone G protein-coupled receptor #233.
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                                                                                                                                                                                               02-JUL-2002
                                                                                                                                                                                                                                                                  AAU95746 standard; protein;
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                               menstrual cycle; appetite sexual motivation; sexual attraction;
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1; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to olfactory and Pheromone G-protein coupled CC receptor (GPCR) or a protein 95% identical to the GPCR, a specific active CC portion and its encoding polymuclectide. Also included are an agonist, CC antagonist or inhibitor of the GPCR or the polymuclectide, a vector and some and comprising the polymucleotide, a cell transformed by the vector, a non-cC comprising the polymucleotide, a cell transformed by the vector, a non-cC compounds which are known or not known to be agonist, antagonists or inhibitors of natural compounds to the GPCR. The receptor-derived cagonists, antagonists, inhibitors or compounds are used as an CC improvement, elimination or substitution of an existing taste and/or a fragrance of (or in) the food and/or cosmetic products. They can also be used in the preparation of medicament in the treatment and/or prevention of amamalian disorder, such as cell migration, sterility, psychotic and neurological disorders, including anxiety, schizophrenia, manic compounds in the preparation for promoting axonal growth, nerve cell connection and nerve regeneration for modulating male and female compression, depression, hormone production and the menstrual cycle, for the prevention or the treatment by stimulation of appetite, sexual attraction, aggression and for promoting or sequence is a human olfactory and pheromone GPCR protein sequence
                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 317; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel pheromone G-protein coupled receptor and receptor-derived agoniant antagonists or inhibitors useful in food or cosmetic products or in the treatment or prevention of neurological disorders such as anxiety and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-330013/36.
N-PSDB; ABK68633.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 710-711; 833pp; English.
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                                                                                                     YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF
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99.7%;
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Pred. No. 2.2e-169;
1; Mismatches 0;
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Best Local S
Matches 317
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Representing sensory perception of one or more odorants identification and design of tastes and odors comprises representative group of n olfactory receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G-coupled olfactory receptor
                                                                                                                                                                                                       Sequence 318 AA;
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13-MAR-2001; 2001US-00804291.
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DB; ABK37540.
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                                           PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120
   MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS 180
                                                                                       MMVDPNGNESSATYFILIGLPGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE
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                                                                                                                                               Conservative
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99.7%;
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                                                                                                                                                         Score 1619; DB 5;
Pred. No. 2.2e-169;
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                                                                                                                                               Mismatches
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Claim 11; Fig

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English

disorders.

The invention relates to a new isolated G protein-coupled receptor

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RESULT 17
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Barber L,
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16-JAN-2001; 2001US-02617765P.
13-JUL-2001; 2001US-0305351P.
17-AUG-2001; 2001US-0313202P.
                                                                                                                                New human G-protein coupled receptor polypeptide, HGPRBMY4, expressed highly in prostate, colon and lung, useful for treating, preventing cancer, immune, neurological, cardiovascular, colon and lung-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2003.
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DB; ACD29000.
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) MINTIER G.
) RAMANATHAN (
) HAWKEN D R.
) CACACE A.
) BARBER L.
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RESULT 18
ABR54523
ID ABR54
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KW Cytos
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Best Local Similarity 99.7
Matches 317; Conservative
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Cytostatic; gene therapy; prostate-specific protein; PSP; human;
                                        Prostate tumour specific protein SEQ ID 920.
                                                                                     28-AUG-2003
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                                                                                                                                                                 standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISPSYLLILKTVLGLTREAQAXAF
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                                                                                                                                                                                                                                                                                                                                                                   GTCVSHVCAVFIFYVPFI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PMY IPLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQMFAIHSLSGMESTVLLA
                                                                                     (first entry)
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Pred. No. 2.2e-169;
1; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to novel prostate-specific proteins (PSP) and their coding sequences. The PSPs and their coding sequences are useful for stimulating an immune response in a patient, or for treating prostate cancer in a patient and for determining, detecting or diagnosing the presence of a cancer in a patient. The present sequence was used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New prostate-specific proteins and genes, useful in gene therapy, particularly for stimulating an immune response in a patient, or prostate cancer in a patient, as well as for diagnosing prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Xu J, Dillon DC, Mitcham JL, Harlocker SL, Jiang Y, Henderson
KaloB MD, Fanger GR, Retter MM, Stolk JA, Day CH, Vedvick TS;
Carter D, Li SX, Wang A, Skeiky YAW, Hepler WT, Hural J;
Mcneill PD, Houghton RL, Vinals Y De BassolsC, Foy TM, Watanabe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-MAY-2001; 2001US-00852911.
29-JUN-2001; 2001US-00895814.
10-DEC-2001; 2001US-00012896.
                                                                                                                                                                                                                                                                                                                                           Sequence 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-MAY-2002; 2002WO-US014753
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; Page 638; 691pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immune response; prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      patient.
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                                                                                                                                                                                                                                       Similarity
                                             GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLFVILANIYLLVPFVLNPIVYGVKTKE
                                                                                MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS
                                                                                                                                                                                                    PMYIFICMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120
IRORILRLFHVATHASBP
                             GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLFVILANIYLLVPFVLNPIVYGVKTKB
                                                                                                                                                                                    PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQMFAIHSLSGMESTVLLA
                                                                                                                                                                                                                                                                                                                                                                        the invention
                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                      99.8%;
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Pred. No. 2.2e-169;
1; Mismatches 0;
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                                                                                                                                                                                                                                                                                           Gaps
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CC The present invention describes antigenic peptides (I) comprising: (a) CC any one of 1601 sequences (see ABP8201) to ABP83619) of 12-24 amino CC acids. Also described: (I) an assay for the detection of a particular G CC protein-coupled receptor (GPCR) or a candidate polypeptide in a sample; and (2) an isolated antibody having high specificity and high affinity or CC avidity for a particular GPCR. (I) can be used as GPCR modulators and in CC gene therapy. The antipodies for GPCRs are useful in detecting an CC antibody against a particular GPCR, and in the production of specific c antibody against a particular GPCR, and in the production of specific CC control of a particular GPCR, and in the production of specific CC resence or absence of corresponding GPCRs. The antigenic peptides for CFCRs and antibodies are useful for detecting the CC presence or absence of corresponding GPCRs. The antigenic peptides for CFCRs and antibodies are useful for diagnosing and designing drugs for treating immune-related diseases, growth-related disease, cell CC regeneration-related diseases, growth-related cell proliferative CC diseases, or autoimmune diseases, e.g. AIDS, Alzheimer's disease, CC atherosclerosis, bacterial, fungal, protozoan or viral infections, CC atherosclerosis, bacterial, fungal, protozoan or viral infections, CC disease, Parkinson's disease, multiple sclerosis, pain, psoriasis, CC anxiety, deports and disease, multiple sclerosis, pain, psoriasis, continuation, schizophrenia, dementia, mental retardation, memory CC loss, epilepsy, asthma, tuberculosis, obesity, nausea, hypertension, CC any other disease, schizophrenia, dementia, mental retardation, memory be used in immunoassays and immunodiagnosis. ABZ42523 to ABZ42869 encode CC exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G protein-coupled receptor; GPCR; antigenic peptide; gene therapy; G protein-coupled receptor modulator; antibody; immune-related disease; growth-related disease; cell regeneration-related disease; AIDS; cancer; immunological-related cell proliferative disease; autoimmune disease; Alzheimer's disease; atherosclerosis; infection; osteoarthritis; allergy; osteoporosis; cardiomyopathy; inflammation; Crohn's disease; diabetes; graft versus host disease; Parkinson's disease; multiple sclerosis; pain; psoriasis; anxiety; depression; schizophrenia; dementia; memory loss; mental retardation; epilepsy; asthma; tuberculosis; obesity; nausea; hypertension; hypotension; renal disorder; rheumatoid arthritis; trauma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 1; 523pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated antigenic peptides e.g., for G protein-coupled receptors (GPCR), useful for diagnosing and designing drugs for treating conditions in which GPCRs are involved, e.g. AIDS, Alzheimer's disease, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-DEC-2000; 2000US-0257144P
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DB; ABZ42822.
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                                             present invention
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23-SEP-1998
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01-AUG-1997;
09-FEB-1998;
25-FEB-1998;
14-JUL-1998;
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2000US-00536857
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                                                                                                                                                                                                                                                                                                                                                   CC The invention relates to an isolated polypeptide comprising no more than CC 11-542 amino acids of ADB13563 comprising a sequence ADB14487. The peptides comprise a fragment ADB13563 of that contain naturally processed CC T-cell epitopes for 3 class I major histocompatibility complex (MHC) CC alleles. ADB13563 is a polypeptide encoded by a human prostate specific CCDNA, one of 648 disclosed as new. Also included are nucleic acids crown, one of 648 disclosed as new. Also included are nucleic acids crown, one of 648 disclosed as new. Also included are nucleic acids crown, one of 648 disclosed as new. Also included are nucleic acids crown, one of 648 disclosed as new. Also included are nucleic acids crown and the proteins and peptides, expression vectors, a host cell crown and the proteins and peptides, expression vectors, a host cell crown appearing a cancer in a patient (comprising contacting a patient CC sample with a binding agent that binds to the peptides or a polypeptide to a predetermined cutcamperising the amount of polypeptide that binds to the appetides or a polypeptide contacting a comparising comparising the peptides or proteins, stimulating or expanding T cells specific for a tumour protein comprising contacting response to the isolated T cell population, treating prostate cancer in a patient with the peptides or antigen presenting chals that express contacting contacting the presence of a cancer in a patient with the peptides or antigen presenting cells that express contacting a patient than a patient. The peptides, nucleic acids encoding, or antigen-presenting cells expressed to respect the populations or attigen protein or expand T cells expressing the nucleic acids encoding, or cancer in a patient. The present sequence is a protein or antigen cells are used to stimulate an immune response or treat comprising cells are used to stimulate an immune response or treat comprised the protein of the printed contact in a patient. The present sequence is a protein the protein contact of the printed contac
                                                                                                                                                                                                                                           Matches
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09-MAY-2001; 2001US-00852911.
29-UUN-2001; 2001US-00895814.
10-DEC-2001; 2001US-00012896.
09-MAY-2002; 2002US-00144678.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polypeptide for use in a vaccine for stimulating an immune response, or for treating or diagnosis cancer, preferably prostate
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US-10-149-316-91
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ALIGNMENTS

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Sequence 2880, Application US/10001469

| Publication No. US20030091562A1 |
| GENERAL INFORMATION: AYA |
| APPLICANT: HAITANO, ARTHUR |
| APPLICANT: APAR, DANIEL |
| APPLICANT: HUBERT, RENE |
| APPLICANT: HUBERT, WARY |
| APPLICANT: PARRS, MARY |
| APPLICANT: PARRS, WARY |
| APPLICANTION NUMBER: US/10/001,469 |
| CURRENT APPLICATION NUMBER: 00/157,902 |
| PRIOR APPLICATION NUMBER: 60/291,118 |
| PRIOR APPLICATION NUMBER: 09/680,728 |
| PRIOR APPLICATION NUMBER: 09/680,728 |
| PRIOR APPLICATION NUMBER: 09/680,728 |
| PRIOR FILING DATE: 2000-10-05 |
| PRIOR APPLICATION NUMBER: 09/680,728 |
| PRIOR FILING DATE: 2000-10-05 |
| PRIOR APPLICATION NUMBER: 09/680,728 |
| PRIOR FILING DATE: 2000-10-05 |
| PRIOR FILING DATE: 2000-10-05 |
| PRIOR FILING DATE: 2000-10-05 |
| PRIOR FILING DATE: 2001-10-05 |

RESULT 2
US-10-147-368-19
; Sequence 19, Application US/10147368
; Publication No. US20030213004A1
; GENERAL INFORMATION:
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ORGANISM: Unknown Organism
PEATURE:
OTHER INFORMATION: Description of Unknown Organism: 101P3A11 encoded amino
OTHER INFORMATION: acid sequence
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Similarity 100.0%; Pred. No. 1.7e-151;
18; Conservative 0; Mismatches 0;
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Sequence 28, Application US/10147368
Publication No. US20030213004A1
GENERAL INFORMATION:
APPLICANT: Agensys, Inc.
APPLICANT: Aya Jakobovits
APPLICANT: Mary Faris
APPLICANT: Arthur B. Raitano
APPLICANT: Robert Kendall Morrison
APPLICANT: Douglas Saffran
APPLICANT: Wangmao Ge

Mary Faris
Arthur B. Raitano
Robert Kendall Morrison
Douglas Saffran
Wangmao Ge

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APPLICANT: Aysunsys, INC.
APPLICANT: Mary Faris
APPLICANT: Mary Faris
APPLICANT: Arthur B. Raitano
APPLICANT: Robert Kendall Morrison
APPLICANT: Robert Kendall Morrison
APPLICANT: Douglas Saffran
APPLICANT: Douglas Saffran
APPLICANT: Pia M. Challita-Eid
TITLE OF INVENTION: NUCLEIC ACIDS AND CORRESPONDING PROTEINS
TITLE OF INVENTION: BRITITLED 10193A11 or PHOR-1 USEFUL IN TREAT
TITLE OF INVENTION: BRITITLED 10193A11 or PHOR-1 USEFUL IN TREAT
TITLE OF INVENTION: BRITITLED 10193A11 or PHOR-1 USEFUL IN TREAT
TITLE OF INVENTION: BRITITLED 10193A11 or PHOR-1 USEFUL IN TREAT
TITLE OF INVENTION NUMBER: US/10/14,368
CURRENT FILING DATE: 2001-12-16
PRIOR APPLICATION NUMBER: US 10/01,469
PRIOR APPLICATION NUMBER: US 10/01,469
PRIOR APPLICATION NUMBER: US 60/291,118
PRIOR APPLICATION NUMBER: US 60/157,902
PRIOR FILING DATE: 2000-10-05
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; TYPE: PRT
; ORGANISM: Homo Sapiens
US-10-147-368-19
RESULT 3
US-10-147-368-28
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LENGTH: 318
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PRIOR APPLICATION NUMBER: US 10/017,066
PRIOR APPLICATION NUMBER: US 10/001,469
PRIOR APPLICATION NUMBER: US 60/291,118
PRIOR PILING DATE: 2001-10-31
PRIOR PILING DATE: 2001-05-15
PRIOR PILING DATE: 2001-05-15
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR PILING DATE: 2000-10-05
PRIOR PILING DATE: 1999-10-05
PRIOR 
Sequence 920, Application US/09759143
Patent No. US20022248A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Henderson, Robert A.
APPLICANT: Kalos, Mitchael D.
APPLICANT: Ranger, Gary R.
APPLICANT: Retter, Marc W.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: USY, Craig H.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Carter, Darrick
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Best Local Similarity
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TITLE OF INVENTION: NUCLEIC ACIDS AND CORRESPONDING PROTEINS
TITLE OF INVENTION: ENTITLED 10193A11 or PHOR-1 USEFUL IN TREATMENT
TITLE OF INVENTION: DETECTION OF CANCER
FILE REFERENCE: 51158-20024.21
CURRENT APPLICATION NUMBER: US/10/147,368
CURRENT FILING DATE: 2002-12-16
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APPLICANT: Wang, Aijun
APPLICANT: Skeiky, Yasiir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C23
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
TYPE: PAT
ORGANISM: Homo sapiens
US-09-759-143-920
APPLICANT: Xu. Jiangchun
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Kalos, Michael D.
APPLICANT: Kalos, Michael D.
APPLICANT: Retter, Marc W.
APPLICANT: Stolk, John A.
APPLICANT: Usy, Craig H.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
APPLICANT: Li, Samuel
APPLICANT: Hang, Aijun
APPLICANT: Hepler, William
APPLICANT: Hepler, William
APPLICANT: Houghton, Raymond L.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C24
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOFTWARE: FRSESEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
TYPE: PRT
ORCANISM: Homo sapiens
US-09-780-669-920
                                                                                                                                                                                                                                                                   ; Sequence 920, Application US/09822827
; Patent No. US20020081680A1
; GENERAL INFORMATION:
    APPLICANT: Xu, Jiangchun
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
; FILE REFERENCE: 210121.534C1
; CURRENT APPLICATION NUMBER: US/09/822,827
; CURRENT FILING DATE: 2001-03-28
; NUMBER OF SEQ ID NOS: 982
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 920
; LENGTH: 318
; TYPE: PRT
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US-09-822-827-920
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)9-822-827-920
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                 MAFDRYVAICHFLRHATVLTLFRVTKIGVAAVVRGAALMAFLFVFIKQLFFCRSNILSHS 180
                                                                                                                     YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF 240
                                                           PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQMFAIHSLSGMESTVLLA
                                                                            PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120
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                                                                                                                                                                              Score 1619; DB 3;
Pred. No. 3.4e-151;
1; Mismatches 0;
                                                                                                                                                                                                           Length 318;
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; Sequence 920, Application US/098; Publication No. US20020192763A1; GENERAL INFORMATION: APPLICANT: Xu. Jiangchun APPLICANT: Dillon, Davin C.; APPLICANT: Mitcham, Jennifer L
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US-09-895-793-920
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; LENGTH: 318
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-886-055-91
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 91, Application US/09886055
Patent No. US20020132273A1
GENERAL INFORMATION:
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APPLICANT: ZOZULYA, SERGEY
TITLE OF INVENTION: RECEPTOR FINGERPRINTING, SENSORY PERCEPTION, AND
TITLE OF INVENTION: BIOSENSORS OF CHEMICAL SENSANTS
FILE REFERENCE: 078003-0277150
CURRENT APPLICATION NUMBER: US/99/886,055
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: 60/213,812
PRIOR PILING DATE: 2000-06-22
RUMBER OF SEQ ID NOS: 522
NUMBER OF SEQ ID NOS: 522
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nes 317; Conserv
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Application US/09895793 o. US20020192763A1

Harlocker,

Susan L.

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APPLICANT: FANGEY, GAIY R.

APPLICANT: FANGEY, GAIY R.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REFERENCE: 210121.534C2

CURRENT APPLICATION NUMBER: US/09/895,793

CURRENT FILING DATE: 2001-06-29

NUMBER OF SEQ ID NOS: 982

SOFTWARE: FRATSEQ for Windows Version 3.0

SEQ ID NO 920

LENGTH: 318

TYPE: PRT

ORGANISM: Homo sapiens
US-09-895-793-920
; Sequence 920, Application US/098958; Publication No. US20020193296A1; GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun; APPLICANT: Mitcham, Jennifer L.; APPLICANT: Harlocker, Susan L.; APPLICANT: Jiang, Yuqiu JaprliCANT: Kalos, Michael D.; APPLICANT: Retter, Marc W.
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Skeiky, Yasir A.W.
Hepler, William T.
Henderson, Robert A.
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Stolk, John A.
Day, Craig H.
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Houghton, Raymond L.
Vinals de Bassols, Carlota
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                                                                                                                             Application US/09895814
o. US20020193296A1
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Kalos, Michael D.
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Pred. No. 3.4e-151;
1; Mismatches 0;
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Sequence 2, Application US/09966459A

Publication No. US20030022237A1

GENERAL INFORMATION:
APPLICANT: FEDER, J.N.
APPLICANT: HAWKEN, D.R.
APPLICANT: HAWKEN, D.R.
APPLICANT: CACACE, A.
APPLICANT: CACACE, A.
APPLICANT: KORNACKER, M.G.
TITLE OF INVENTION: A NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGP!
TITLE OF INVENTION: EXPRESSED HIGHLY IN PROSTATE, COLON, AND LUNWELLE REFERENCE: D0039NP
CURRENT APPLICATION NUMBER: US/09/966,459A
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C26
CURRENT APPLICATION NUMBER: US/09/895,814
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 990
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
TYPE: PRT
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US-09-895-814-920
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Best Local S
Matches 317
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APPLICANT:
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Similarity 99.7%;
17; Conservative
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Houghton, Raymond L.
Vinals de Bassols, Carlota
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Hepler, William T.
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Vedvick, Thomas
Carter, Darrick
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Thomas
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Pred. No. 3.4e-151;
1; Mismatches 0;
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TITLE OF INVENTION. HUMAN OLFACTORY RECEPTORS AND GENES ENCODING SAME FILE REFERENCE: P 0278005
CURRENT APPLICATION NUMBER: US/09/804,291
CURRENT PILING DATE: 2001-03-13
PRIOR FILING DATE: 2000-03-13
PRIOR FILING DATE: 2000-03-24
PRIOR PILING DATE: 2000-03-24
PRIOR PILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/199,335
PRIOR APPLICATION NUMBER: 60/199,335
PRIOR APPLICATION NUMBER: 60/213,349
PRIOR PILING DATE: 2000-04-26
PRIOR APPLICATION NUMBER: 60/207,702
PRIOR APPLICATION NUMBER: 60/207,702
PRIOR APPLICATION NUMBER: 60/207,702
PRIOR PILING DATE: 2000-06-26
PRIOR APPLICATION NUMBER: 60/213,849
PRIOR APPLICATION NUMBER: 60/226,534
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; TYPE: PRT
; ORGANISM: Homo s
US-09-966-459A-2
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PRIOR APPLICATION NUMBER: 60/235,833
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/261,776
PRIOR FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: 60/305,351
PRIOR FILING DATE: 2001-07-13
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PRIOR FILING DATE: 2001-08-17
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 91, Application US/09804291 Publication No. US20030088059A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 317; Conserv
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OR APPLICATION NUMBER: 60/207,702
OR FILING DATE: 2000-05-26
OR APPLICATION NUMBER: 60/213,849
OR APPLICATION NUMBER: 60/213,849
OR APPLICATION NUMBER: 60/236,534
OR APPLICATION NUMBER: 60/230,732
OR APPLICATION NUMBER: 60/230,732
OR FILING DATE: 2000-09-07
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Pred. No. 3.4e-151;
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PRIOR FILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 529
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 91
LENGTH: 318
TYPE: PRT
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US-09-804-291-91
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Best Local Similarity
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APPLICANT: Fanger, Gary R.
APPLICANT: Wantanabe, Yoshihiro
APPLICANT: Wantanabe, Yoshihiro
APPLICANT: Meagher, Medeleine Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C27
CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
                                                                                                                              APPLICANT:
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APPLICANT: Dillon, Davi
APPLICANT: Mitcham, Jen
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Stolk, John A.
Day, Craig H.
Vedvick, Thomas 9
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Vedvick, Thomas
Carter, Darrick
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Mitcham, Jennifer L.
                                                                                                                                             McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
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Hepler, William T.
                                                                                                                                                                                                                                                                                                                                                                     Jiang, Yuqiu
Kalos, Michael D.
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                                                                                                                               Foy, Teresa
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5. US20020183251A1
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Pred. No. 3.4e
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CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/257,144
PRIOR FILING DATE: 2000-12-19
INVMEER OF SEQ ID NOS: 2292
SOFTWARE: Patentin version 3.1
SEQ ID NO 434
ILENGTH: 318
TYPE: PRT
ORGANISM: Homo sapiens
US-10-225-567A-434
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US-10-225-567A-434
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Best Local (
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Publication No. US20030113798A1
GENERAL INFORMATION:
APPLICANT: LifeSpan Biosciences
APPLICANT: Brown, Joseph P.
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Best Local Similarity 99.7
Matches 317; Conservative
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                                                                                                                                                                                             Matches
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APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
TITLE OF INVENTION: ANTICENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 1920-4-4
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Pred. No. 3.4e-151;
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APPLICANT: ASAI, KIYOSHI
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR PILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2430
NUMBER OF SEQ ID NOS: 2430
RESULT 15
US-10-144-678A-920
US-10-144-678A-920
; Sequence 920, Application US/10144678A
; Publication No. US20030157089A1
; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
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US-10-017-161-868
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Best Local (
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TYPE: PRT
ORGANISM: Homo sapiens
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5. US20030143668A1
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99.7%; Pred. No. 3.4e-151;
tive 1; Mismatches 0;
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APPLICANT:

Kalos, Michael D.

Susan L. iu Robert A.

Yuqiu

Mitcham, Jennifer

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APPLICANT:

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APPLICANT: Watanabo, Yoshihiro

APPLICANT: Deng, Ta

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REFERENCE: 21011.427C28

CURRENT APPLICATION NUMBER: US/10/144,678A

CURRENT FILING DATE: 2002-08-12

NUMBER OF SEQ ID NOS: 1033

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 920

LENGTH: 318

TYPE: PRT

ORGANISM: Homo sapiens

US-10-144-678A-920
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US-10-294-025-920
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Sequence 920, Application US/10294025
Publication No. US20030185830A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Kalos, Michael D.
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Best Local (
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                                                                                                                                                        IRQRILRLFHVATHASEP 318
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Houghton, Raymond L.
Vinals y de Bassols, Carlota
Foy, Teresa M.
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Hepler, William T.
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Vedvick, Thomas S.
Carter, Darrick
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99.7%;
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Pred. No. 3.4e-151;
1; Mismatches 0;
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APPLICANT: Wernet, Kumud
APPLICANT: Wernet, Corine
APPLICANT: Casman, Stacle J
APPLICANT: Casman, Stacle J
APPLICANT: Casman, Stacle J
APPLICANT: Wolenc, Adam R
APPLICANT: Boderna, Steven K
APPLICANT: Hishnu, Vishun S
APPLICANT: Hishnu, Vishun S
APPLICANT: Spytek, Kimberly A
APPLICANT: Syytek, Kimberly A
APPLICANT: Gusev, Valdimir
APPLICANT: Gusev, Vladimir
TITLE OF INVENTION: NO. US20030195335A1el Proteins and Nucleic Acid
FILE REFERENCE: 15966-748
CURRENT APPLICATION NUMBER: 00/103/664
FRIOR FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: 60/193/664
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: 60/194/614
PRIOR APPLICATION NUMBER: 60/194/614
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: 60/195/063
PRIOR APPLICATION NUMBER: 60/195/066
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: 60/195/067
PRIOR APPLICATION NUMBER: 60/195/067
PRIOR APPLICATION NUMBER: 60/195/067
PRIOR APPLICATION NUMBER: 60/195/067
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FILE REFERENCE: 210121.427C29
; CURRENT APPLICATION NUMBER: US/10/294,025
; CURRENT FILING DATE: 2002-11-12
; NUMBER OF SEQ ID NOS: 1038
; SOFTMARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 920
; SEQ ID
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US-10-044-643-26
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Best Local Similarity 99.7
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Pred. No. 3.4e-151;
1; Mismatches 0;
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APPLICATION NUMBER: 60/195,068 FILING DATE: 2000-04-06

APPLICATION NUMBER: 60/195,069

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RESULT 18
US-10-147-368-32
US-10-147-368-32
; Sequence 32, Application US/10147368
; Publication to US20030213004A1
; GENERAL INFORMATION:
; APPLICANT: Agensys, Inc.
; APPLICANT: Aya Jakobovits
; APPLICANT: Mary Faris
; APPLICANT: Arthur B. Raitano
**PPLICANT: Robert Kendall Morrison
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NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 26
LENGTH: 318
TYPE: PRT
ORGANISM: Homo sapiens
US-10-044-643-26
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Best Local Similarity 99.7%;
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FILING DATE: 2000-07-27
APPLICATION NUMBER: 60/221,325
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FILING DATE: 2001-01-23
APPLICATION NUMBER: 60/263,433
FILING DATE: 2001-01-23
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APPLICATION NUMBER: 60/224,588
FILING DATE: 2000-08-11
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FILING DATE: 2000-04-06
APPLICATION NUMBER: 60/219,855
FILING DATE: 2000-07-21
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1 MYVDPNGNESSATYFILIGLPGLEBAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE
1 MYVDPNGNESSATYFILIGLPGLEBAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE
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Pred. No. 3.4e-151;
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APPLICANT: Douglas Saffran

APPLICANT: Wangmao Ge

APPLICANT: Wangmao Ge

TITLE OF INVENTION: NUCLEIC ACIDS AND CORRESPONDING PROTEINS
TITLE OF INVENTION: NUCLEIC ACIDS AND CORRESPONDING PROTEINS
TITLE OF INVENTION: DETECTION OF CANCER
FILE REFERENCE: 51158-20024.21
CURRENT PAPLICATION NUMBER: US/10/147,368
CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: US 10/017,066
PRIOR APPLICATION NUMBER: US 10/01,469
PRIOR FILING DATE: 2001-10-31
PRIOR APPLICATION NUMBER: US 60/291,118
PRIOR APPLICATION NUMBER: US 60/291,118
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR PILING DATE: 2000-10-05
PRIOR PILING DATE: 1999-10-05
PRIOR FILING DATE: 1999-10-05
APPLICANT: HAGA, TATSUYA
ITITLE OF INVENTION: NOVEL G-PROTEIN COUPLED RE
FILE REFERENCE: 31671-186347
CURRENT APPLICATION NUMBER: US/10/343,650A
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: J2000/237818
PRIOR FILING DATE: 2000-08-04
PRIOR APPLICATION NUMBER: JP 2001/34434
PRIOR FILING DATE: 2001-02-13
NUMBER OF SEQ ID NOS: 694
SOFTWARE: PALEENTIN Ver. 2.1
SEQ ID NO 158
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US-10-343-650A-158
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LENGTH: 318
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Best Local Similarity 99.7%;
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TYPB: PRT
ORGANISM: Homo 9
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                                                                                                                                                                                                                                                                                                        Application US/10343650A to. US20040067499A1
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Pred. No. 3.4e-151;
1; Mismatches 0;
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Sequence 10, Application US/10692605

Publication No. US20040091928A1

GENERAL INFORMATION:

APPLICANT: VOGeli, Gabriel

APPLICANT: VOGeli, Gabriel

APPLICANT: Worklant, Kalpana

ITILE OF INVENTION: G PROTEIN-COUPLED RECEPTORS EXPRESSED IN BRAIN

FILE REFERENCE: 28341/6276.NX1

CURRENT APPLICATION NUMBER: US/10/692,605

CURRENT FILING DATE: 2000-01-0-27

PRIOR APPLICATION NUMBER: US 09/484,794

PRIOR APPLICATION NUMBER: US 09/484,799

PRIOR APPLICATION NUMBER: US 09/454,399

PRIOR APPLICATION NUMBER: US 09/454,399

PRIOR FILING DATE: 1999-10-28

PRIOR APPLICATION NUMBER: US 09/429,517

PRIOR APPLICATION NUMBER: US 09/429,676

PRIOR APPLICATION NUMBER: US 09/429,676

PRIOR APPLICATION NUMBER: US 09/429,676

PRIOR FILING DATE: 1999-10-28

PRIOR APPLICATION NUMBER: US 09/429,695

PRIOR FILING DATE: 1999-10-28

PRIOR APPLICATION NUMBER: US 09/428,114

PRIOR APPLICATION NUMBER: US 09/428,095

PRIOR APPLICATION NUMBER: US 09/428,090

PRIOR FILING DATE: 1999-10-27

PRIOR PILING DATE: 1999-10-27
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US-10-692-605-10
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-343-650A-158
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                                                    Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 75
SOPTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 318
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Best Local Similarity 99.7%;
Matches 317; Conservative
TYPE: PRT
ORGANISM: Homo sapiens
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US-11-190-188-17
US-10-055-877-23
US-10-055-877-23
US-11-190-188-13
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US-10-055-877-234
US-10-055-877-234
US-10-055-877-234
US-10-055-877-234
US-10-051-538-18
US-11-190-188-11
US-10-051-538-19
US-11-170-153-2
US-11-170-153-
  sequence
seq
  1324, App 119, App 11
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243

246 183 187

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APPLICANT: Skelky, Yasir A.

TITLE OF INVENTION: PROSTATE-SPECIFIC POLYPEPTIDES AND FUSION
TITLE OF INVENTION: PROSTATE-SPECIFIC POLYPEPTIDES AND FUSION
TITLE OF INVENTION: POLYPEPTIDES THEREOF
FILE REFERENCE: 210121.427C31
CURRENT APPLICATION NUMBER: US/11/234,786
CURRENT FILING DATE: 2005-09-23
FRIOR APPLICATION NUMBER: US 09/568,857
PRIOR FILING DATE: 2000-05-09
PRIOR FILING DATE: 2000-05-07
PRIOR PRIOR DATE: 2000-05-17
PRIOR APPLICATION NUMBER: US 09/536,857
PRIOR PRILING DATE: 2000-05-114
                                                                                                                                                                                                                                                                                  NUMBER FAST:
SOFTWARE: FAST:
SEQ ID NO 527
FUNCTH: 320
                                                                                                                                  Query Match
Best Local Similarity 59...
182; Conservative
                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Homo sapiens
:-11-234-786-527
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Sequence 527, Application US/11234786
Publication No. US20060024301A1
GENERAL INFORMATION:
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FILING DATE: 1999-04-09
ALPLICATION NUMBER: US 09/232,149
FILING DATE: 1999-01-15
                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 1998-09-23
APPLICATION NUMBER: US 09/115,453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 1999-11-12
APPLICATION NUMBER: US 09/352,616
FILING DATE: 1999-07-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 09/439,313
                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 1998-02-25
                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 1998-07-14
APPLICATION NUMBER: US 09/030,607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 09/159,812
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                                                                        MLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYV 127
    MLAAIDLALSTSTMPKILALFWFDSREISFEACLTQMFFIHALSAIESTILLAMAFDRYV
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Vedvick, Thomas S.
Carter, Darrick
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Reed, Steven G.
Kalos, Michael D.
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Mitcham, Jennifer L.
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US-10-511-538-172
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Pred. No. 3e-82;
S5; Mismatches 67
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                                                                                                                                                      67,
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Sequence 172,
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    123
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                                              APPLICANT: Casman, Stacie
APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-39
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR PILING DATE: 2001-01-26
PRIOR PILING DATE: 2001-01-26
PRIOR PILING DATE: 2001-01-30
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APPLICATION NUMBER: 60/272,870 FILING DATE: 2001-03-02 APPLICATION NUMBER: 60/275,990
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Miller, Charles
Miller, Velizar
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Spytek, Kimberly
Ratelli, Luca
Kekuda, Ramesh
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Zerhusen, Bryan
Andrew, David
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Baumgartner, Jason
Shimkets, Richard
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Taupier Jr., Raymond
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Burgess, Cahterine
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o. US20050288241A1
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; PRIOR FILING DATE: 2001-03-14
; PRIOR APPLICATION NUMBER: 60/275,927
; PRIOR FILING DATE: 2001-03-14
; Remaining Prior Application data removed - Sei NUMBER OF SEQ ID NOS: 512
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 336
; ERNOTH: 329
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-336
                                                                                                                                                                                                                                                                                                                                                                     US-10-055-877-339

; Sequence 339, Application US/10055877
; Publication No. US20050288241A1
; GENERAL INFORMATION:
; APPLICANT: DeCristofaro, Marc
; APPLICANT: Padigaru, Muralidhara
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6 NGNESSATYFILIGLEGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIF
Wolenc, Adam
Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
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Zerhusen, Bryan
Andrew, David
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Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
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Miller, Charles
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Burgess, Cahterine
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Kekuda, Ramesh
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                                                                                                                  Bisen, Andrew
                                                                                                                                                                Mezes, Peter
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Pred. No. 2.5e-74;
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; ORGANISM: Homo sapiens
US-10-055-877-339
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PRIOR PILLING DATE: 2001-01-23
PRIOR PILLING DATE: 2001-01-24
PRIOR PILLING DATE: 2001-01-24
PRIOR PILLING DATE: 2001-01-24
PRIOR PILLING DATE: 2001-01-25
PRIOR PILLING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR PILLING DATE: 2001-01-25
PRIOR PILLING DATE: 2001-01-25
PRIOR PILLING DATE: 2001-01-25
PRIOR PILLING DATE: 2001-01-30
PRIOR PILLING DATE: 2001-01-30
PRIOR PILLING DATE: 2001-03-14
PRIOR PILLING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR PILLING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR PILLING DATE: 2001-03-14
PRIOR PILLING DATE: 2001-03-14
PRIOR PILLING DATE: 2001-03-14
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NUMBER OF SEQ ID NOS: 512
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 339
LENGTH: 312
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Best Local (
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APPLICANT: Boldog, Ference
TITLE OP INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
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CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
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                                                                                      245
                                                                                                                     247 VCAVFIFYVDFIGLSMVHRFSKRRDSPLPVILANIYLLVDFVLNDIVYGVKTKEIRQRIL 306
                                                                                                                                                                                                              188 MKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLG-LTREAQAKAFGTCVSH 246
                                      307 RLFH 310
                                                                                                                                                                                                                                                                          125 AICDPLRYATVLTTEVIAAMGLGAAARSFITLFPLPFLIKKLPICRSNVLSHSYCLHPDM
                                                                                                                                                                                                                                                                                                  128 AICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQDV 187
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RMFH 308
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                                                                                      ILAVLAFYVPMIGVSTVHRFGKHVPCYIHVLMSNVYLFVPPVLNPLIYSAKTKEIRRAIF
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Pred. No. 3.7e-72;
s1; Mismatches 89;
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; Sequence 8, Application US/11190188; Publication No. US20060035331A1; GENERAL INFORMATION:
; APPLICANT: INCYTE GENOMICS, INC.; APPLICANT: LAL, Preeti; APPLICANT: TANG, Y. Tom
; APPLICANT: PATTERSON, Chandra

LAL, Preeti TANG, Y. Tom PATTERSON, Chandra

RESULT 4 US-11-190-188-8

Monique G.

Leo L.

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RESULT 5
US-10-511-538-70
VS-10-511-538-70
Sequence 70, Application US/10511538
Publication No. US20060026700A1
GENERAL INFORMATION:
APPLICANT: OriGene Technologies, Inc
TITLE OF INVENTION: TISSUE SPECIFIC GENES AND FILE REFERENCE: 16U 200 PCT
CURRENT APPLICATION NUMBER: US/10/511,538
CURRENT FILING DATE: 2004-10-18
PRIOR APPLICATION NUMBER: US 60/372,669
PRIOR FILING DATE: 2002-04-16
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PRIOR FILING DATE: 2000-03-03; 2000-03-10; 2000-03-17; 2000-03-20
NUMBER OF SEQ ID NOS: 42
SOPTWARE: PERL Program
SEQ ID NO 8
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Best Local Similarity 49.8%; Pred. No. 2.1e-68;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAM 121
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AU-YOUNG, Janice
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; TYPE: PRT
; ORGANISM: Homo &
US-10-511-538-70
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Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
Ratelli, Luca
                                      Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Padigaru, Muralidhara
Miller, Charles
                                                                                                                                                          Mezes, Peter
Patturajan, Meera
Burgess, Cahterine
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Zerhusen, Bryan
Andrew, David
               Vernet,
                                                                                                            Wolenc, Adam
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o. US20050288241A1
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Query Match 50.1%; Score 812.5; DB 6; Best Local Similarity 50.7%; Pred. No. 4.9e-67; Matches 151; Conservative 59; Mismatches 87;
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OR APPLICATION NUMBER: US 60/424,336
OR FILING DATE: 2002-11-07
OR APPLICATION NUMBER: US 60/374,823
OR FILING DATE: 2002-04-24
OR APPLICATION NUMBER: US 60/376,558
OR FILING DATE: 2002-05-01
OR APPLICATION NUMBER: US 60/381,366
OR APPLICATION NUMBER: US 60/403,648
OR FILING DATE: 2002-08-08
OR FILING DATE: 2002-08-16
OR FILING DATE: 2002-08-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                12 TAFLLVGIPGLEHLHIWISIPFCLAYTLALLGNCTLLLIIQADAALHEPWYLFLAWLAAI
                       IFYVPFIGLSMVHRFSKRRDSPLFVILANIYLLVPFVLNFIVYGVKTKEIRQRILRLF 309
                                                                                                         GDTSFNNIYGIAVAMFIVVLDLLLVILSYIFILQAVLLLASQEAHYKAFGTCVSHIGAIL 251
                                                                                                                                                                                                                                                                                                                                                                 DILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYVAICHP 132
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APYTTVVISSVMHRVARHAAPHVHILLANFYLLFPPMVNPIIYGVKTKQIRESILGVF 309
                                                                                                                                               DDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSHVCAVF 251
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RESULT 7
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F.190-188-10
F. Sequence 10, Application US/11190188
Publication No. US20060035331A1
GENERAL INFORMATION:
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US-10-055-877-335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 512
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 335
LENGTH: 342
TYPE: PRT
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Best Local Similarity
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PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
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TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids
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OR FILING DATE: 2001-01-23
OR APPLICATION NUMBER: 60/263,799
OR FILING DATE: 2001-01-24
OR APPLICATION NUMBER: 60/264,117
OR FILING DATE: 2001-01-25
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APPLICATION NUMBER: 60/275,990
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FILING DATE: 2001-01-25
APPLICATION NUMBER: 60/264,478
FILING DATE: 2001-01-26
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                                                                                                                                                        DVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCV 244
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46.7%; Pred. No. 8.4e-66;
ative 72; Mismatches 89;
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; Sequence 338, Application US/100
; Publication No. US20050288241A1
; GENERAL INFORMATION;
; APPLICANT: DeCristofaro, Marc
; APPLICANT: Paddgaru, Muralidha
; APPLICANT: Miller, Charles
; APPLICANT: Tchernev, Velizar
; APPLICANT: Zhong, Mei

Application US/10055877 o. US20050288241A1

Padigaru, Muralidhara Miller, Charles Tchernev, Velizar

APPLICANT:

Anderson, David Ballinger, Robert

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RESULT 8
US-10-055-877-338
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FILE REFERENCE: PI-0044 PCT
CURRENT APPLICATION NUMBER: US/11/190,188
CURRENT FILING DATE: 2005-07-27
PRIOR APPLICATION NUMBER: US/10/220,382
PRIOR FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: 60/186,854; 60/188,384; 60/190,453; 60/190,730
PRIOR FILING DATE: 2000-03-03; 2000-03-10; 2000-03-17; 2000-03-20
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PERL Program
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 47.5%; Score 771; DB 7; Length Best Local Similarity 47.3%; Pred. No. 3.1e-63; Matches 140; Conservative 71; Mismatches 83; Indel
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ORGANISM: Homo sapiens
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                                                                          YIPSVFSFLTHRFGHQIPGYIHILVANLYLIIPPSLNPIIYGVRTKQIRERVLYVF 308
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WALSH, Roderick T.
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TRIBOULEY, Catherine
LU, Dyung Aina M.
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TANG, Y. To
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APPLICANT:

Gerlach, Valerie Spytek, Kimberly

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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR PPLICATION NUMBER: 60/263,598
PRIOR PPLICATION NUMBER: 60/263,598
PRIOR PILING DATE: 2001-01-33
PRIOR PILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR PILING DATE: 2001-01-24
PRIOR PILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
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US-10-055-877-338
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SOPTWARE: Pate
; SEQ ID NO 338
; SURTH: 321
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Best Local Similarity 45.2%;
Matches 140; Conservative 73
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TITLE OF INVENTION: Novel Polypeptides and Nucleic
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APPLICATION NUMBER: 60/275,990
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YCLHQDVMKLACDDIRVNYVYGLIVIISAIGLDSLLISESYLLILKTVLGL-TREAQAKA 239
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                                                         MAPDRYIAICNPLRYSSILTNSRIIKIGLTIIGRSFFFFITPPIICLKFFNYCHFHILSHS 186
                                                                                                                                     PMPYPLSMLALTDLCMGLSTVYTVLGILWRIIREISLDSCIAQSYPIHGLSPMESSVLLT 126
                                                                                                                                                                       PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120
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Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
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Taupier Jr., Raymond
Pena, Carol
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Burgess, Cahterine
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Mezes, Peter
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Zerhusen, Bryan
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Pred. No. 4.4e-62;
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; LENGTH: 356
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No: 7472363CD1
US-11-190-188-9
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US-11-190-188-9
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45.7%; Score /**, ---
Best Local Similarity 46.7%; Pred. No. 1.6e-6
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CURRENT APPLICATION NUMBER: US/11/190,188
CURRENT FILING DATE: 2005-07-27
PRIOR APPLICATION NUMBER: US/10/220,382
PRIOR FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: 60/186,854; 60/188,384; 60/190,453; 60/190,730
PRIOR FILING DATE: 2000-03-03; 2000-03-10; 2000-03-17; 2000-03-20
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PERL Program
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 261
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                                                                                                                                                                                   82 PKMLAIFWENSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYVAICHPLRHATVLTL 141
                                                                                                                                                                                                                      99
                                                                                                                                                                                                                                    T: BOROWSKY, Mark L.
INVENTION: G-PROTEIN COUPLED RECEPTORS
                                                         GLIVIISAIGLDSLLISESYLLILKTVLGL-TREAQAKAFGTCVSHVCAVFIFYVPFIGL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EIRQRILRLF 309
SMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKBIRQRILRLF 309
                               GLEV-VSEFVLNLVLIGISYVYILRAVERLPSHDAQLKALSTCGAHVGVICVFVIPSVFS
                                                                                                                                                       GLEDFHMWISGPFCSVYLVALLGNATILLVIKVEQTLREPMFYFLAILSTIDLALSATSV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            POTCISHICAVLVEYIPIISLTMVHREGKHLSPVAHVLIGNIYILFPPLMNPIIYSVKTO 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FGTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTK 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TANG, Y. Tom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WALSH, Roderick T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HARLAND,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AU-YOUNG, Janice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SHIH, Leo L.
TRIBOULEY, Cather
LU, Dyung Aina M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Terence P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Junming
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Catherine
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                                                                                                                                                                                                                                                                                                  Score 742; DB 7;
Pred. No. 1.6e-60;
                                                                                                                                                                                                                                                                                    82; Indels
                                                                                                                                                                                                                                                                                                               Length 356;
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                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                            185
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                                                                                                                            201
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CURRENT APPLICATION NUMBER: US/11/124,367A
CURRENT FILING DATE: 2005-05-09
PRIOR APPLICATION NUMBER: US 60/568,846
PRIOR FILING DATE: 2004-05-07
PRIOR APPLICATION NUMBER: US 60/582,609
PRIOR PILING DATE: 2004-06-25
PRIOR APPLICATION NUMBER: US 60/599,554
PRIOR APPLICATION NUMBER: US 60/599,554
PRIOR PILING DATE: 2004-08-09
NUMBER OF SEQ ID NOS: 34460
SOPTWARE: PASTSEQ for Windows Version 4.0
SEQ ID NO 496
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; ORGANISM: Homo sapiens
US-11-124-367A-496
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US-11-124-367A-496
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                                                                                                                                                               Sequence 514, Application US/11124367A Publication No. US20060024700A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 496, Application US/11124367A Publication No. US20060024700A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches 135;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
APPLICANT: Michele Cargill
APPLICANT: Hongjin Huang
TITLE OF INVENTION: Genetic Polymorphisms Associated with
TITLE OF INVENTION: Fibrosis Methods of Detection and Uses
FILE REFERENCE: CL001519.ORD
CURRENT APPLICATION NUMBER: US/11/124,367A
CURRENT FILING DATE: 2005-05-09
PRIOR APPLICATION NUMBER: US 60/568,846
PRIOR FILING DATE: 2004-05-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Michele Cargill
APPLICANT: Hongjin Huang
TITLE OP INVENTION: Genetic Polymorphisms Associated with
TITLE OP INVENTION: Fibrosis Methods of Detection and Uses Thereof
FILE REFERENCE: CL001519.ORD
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                                                                                                                                                                                                                                                                                                                                   306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62 AMLAATDLGVTLTIMPTVLGVLWLDHREIGHGACFSQAYFIHTLSVMESGVLLAMAYDCF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ITHRNPLRYTSILTNTQVMKIGVRVLTRAGLSIMPIVVRLHWPPYCRSHVLSHAFCLHQD
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Pred. No. 3.6e-60;
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APPLICANT: DeCristofaro, Marc
APPLICANT: Dedigaru, Muralidha;

APPLICANT: Miller, Charles

APPLICANT: Tchernev, Velizar

APPLICANT: Zhong, Mei

APPLICANT: Anderson, David

APPLICANT: Ballinger, Robert

APPLICANT: Gerlach, Valerie

APPLICANT: Spytek, Kimberly

APPLICANT: Ratelli, Luca

APPLICANT: Ratelli, Luca

APPLICANT: Kekuda, Ramesh

APPLICANT: Kekuda, Ramesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; PRIOR APPLICATION NUMBER: US 60/582,609; PRIOR FILING DATE: 2004-06-25; PRIOR APPLICATION NUMBER: US 60/599,554; PRIOR FILING DATE: 2004-08-09; NUMBER OF SEQ ID NOS: 34460; SOPTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-055-877-123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 123, Appublication No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 45.4%;
Best Local Similarity 45.8%;
                                                     APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
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APPLICANT:
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                     APPLICANT
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                                                                       Shimkets, Richard
Gusev, Vladimir
Vernet, Corine
                                                                                                                                                                                                                     Andrew, David
Mezes, Peter
                                                                                                                                                                                                                                                         Guo, Xiaojia
Zerhusen, Bryan
   Shenoy, Suresh
Li, Li
                                                                                                                         Wolenc, Adam
Baumgartner, Jason
                                                                                                                                                                 Burgess, Cahte
Bisen, Andrew
                                    Pena,
                                                   Taupier Jr., Raymond
                                                                                                                                                                                  Patturajan, Meera
Burgess, Cahterine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/10055877 b. US20050288241A1
                                    Carol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Muralidhara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  68; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 736.5; DB 7
Pred. No. 4.4e-60;
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241 SHICCVLVFYVTVIGLSLIHRFGKQVPHIVHLIMSYAYFLFPPLMNPITYSVKTKQIQNA 300
                                                                                                                                                                                                                                                       186 DVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCV
                                                                                                                                                                                                                                                                                                                  128 AICHPLRHATVLTLPRVTKIGVAAVVRGAALMAP--LPVFIKQLPFCRSNILSHSYCLHQ
                                                                                                                                                                                                                                                                                                                                                                                 63 MLAATDLGLALTIMPTVLGVLWLDHREIGSAACFSQAYFIHSLSFLESGILLAMAYDRFI
                                                                                                                                                                                                                                                                                                                                                                                                                                                68 MLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYV 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 SSSSHPFLLTGFPGLEEAHHWISVFFLFMYISILFGNGTLLLLIKEDHNLHEPMYFFLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8 NESSATYFILIGLEGLEBAQFWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHBEMYIFLC
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                                                                                                                             DVIKLACADTTENRLYPAVLVVFIFVLDYLIIFISYVLILKTVLSIASREERAKALITCV
                                       SHVCAVEIFYVEFIGLSMYHRFSKRRDSPLFVILANIYLLVFPVLNFIVYGVKTKEIRQR 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 7;
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244 180 185 122

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US-10-055-877-337; Sequence 337, Application US/10055877; Publication No. US20050288241A1
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                                                                                                                                                                                                         RESULT 13
                                                        GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Tcherney, Velizar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
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SEQ ID NO 123
LENGTH: 303
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 APPLICANT:
APPLICANT:
APPLICANT:
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ORGANISM: Homo sapiens
-10-055-877-123
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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/272,870 FILING DATE: 2001-03-02 APPLICATION NUMBER: 60/275,990 FILING DATE: 2001-03-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/264,478
FILING DATE: 2001-01-26
APPLICATION NUMBER: 60/263,351
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/275,927 FILING DATE: 2001-03-14
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                                                                                                                                                                                                                                                                                                                              185
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                                                                                                                                                                                                                                                       YIHMLSLSLVYRYGRSAPRVVHSVMANVYLLLPPVLNPIIDSVKTKQIRKAMLSL 299
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                                                                                                                                                                                                                                                                            YVPFIGLSMYHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKBIRQRILRL 308
                                                                                                                                                                                                                                                                                                                                                                                                                              HATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQDVMKLACDD 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYVAICHPLR 134
                                                                                                                                                                                                                                                                                                                              IRANSICGLTDLILTTGIDTPCIVLSYILIIHSVLRIASPBEWHKVFSTCVSHVGAVAFF
                                                                                                                                                                                                                                                                                                                                                              IRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSHVCAVFIF 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FLITGIPGLESAHVWISIPFCCFYAIALSGNSVILFVIITQQSLHEPMYYFLFRLSATDL
Anderson, David
Ballinger, Robert
Gerlach, Valerie
                                                 Zhong, Mei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Casman, Stacie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       44.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 726.5; DB 6
Pred. No. 3.5e-59;
8; Mismatches 98
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APPLICANT: Shenoy, Suresh
APPLICANT: Shenoy, Suresh
APPLICANT: Shenoy, Suresh
APPLICANT: Li, Li
APPLICANT: Lasman, Stacie
APPLICANT: Casman, Stacie
APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
CURRENT PAPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/264,117
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Homo sapiens US-10-055-877-337
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NUMBER OF SEQ ID NOS: 512
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 337
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Best Local (
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PRIOR APPLICATION NUMBER: 60/264,478
PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 32
TYPE: PRT
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APPLICATION NUMBER: 60/275,990
FILING DATE: 2001-03-14
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                                         186 DVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCV 244
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Similarity 46.4%;
41; Conservative 5
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EVIKYTYSKPWISSFWGLFLQLYLNGTDVLFILFSYVLILRTVLGIVARKKQQKALSTCV
                                                                                          FVAICNPLNYATILTDRMVLVIGLVICIRPAVFLLPLLVAINTVSFHGGHELSHPFCYHP
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Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
Vernet, Corine
Taupier Jr., Raymond
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Mezes, Peter
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Burgess, Cahterine
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Zerhusen, Bryan
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Pred. No. 2.1e-58;
8; Mismatches 104;
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RESULT 15
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; ORGANISM: Homo sapiens
US-10-511-538-225
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US-10-511-538-225
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SEQ ID NO 225
LENGTH: 291
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Best Local :
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PRIOR
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TITLE OF INVENTION: TISSUE SPECIFIC GENES AND
FILE REFERENCE: 16U 200 PCT
CURRENT APPLICATION NUMBER: US/10/511,538
CURRENT FILING DATE: 2004-10-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/411,882 PRIOR FILING DATE: 2002-09-20
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PRIOR FILING DATE: 2002-04-16
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Local Similarity 41.3%; Pred. No. 5.1e-52;
hes 126; Conservative 59; Mismatches 98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 60/376,558
FILING DATE: 2002-05-01
APPLICATION NUMBER: US 60/381,366
FILING DATE: 2002-05-20
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APPLICATION NUMBER: US 60/374,823
FILING DATE: 2002-04-24
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                                                                                        282
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                                                                                                                                 ILRLF 309
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                                                                                        ILLLF 286
                                                                                                                                                                        SHVCVILIFYIPAFFSILAHCFGHNVPHTFHIMFANLYVIIPPALNSIVYRIKTKQIQNR
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; Publication No. US20060026700A1
; GENERAL INFORMATION:
; APPLICANT: Origene Technologies, Inc.
; TITLE OF INVENTION: TISSUE SPECIFIC GENES AN
; FILE REFERENCE: 16U 200 PCT
; CURRENT FILING DATE: 2004-10-18
; DRICH ADDITORATION NUMBER: US/10/511,538
; DRICH ADDITORATION NUMBER: US/10/511,538
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PRIOR FILLING DATE: 2002-04-16
PRIOR PELLORION NUMBER: US 60/411,882
PRIOR FILLING DATE: 2002-09-20
PRIOR FILLING DATE: 2002-09-20
PRIOR PELLORITON NUMBER: US 60/424,336
PRIOR PELLORITON NUMBER: US 60/374,823
PRIOR PELLORITON NUMBER: US 60/374,823
PRIOR PELLORITON NUMBER: US 60/374,823
PRIOR PELLORITON NUMBER: US 60/376,558
PRIOR APPLICATION NUMBER: US 60/376,558
PRIOR APPLICATION NUMBER: US 60/381,366
PRIOR FILLING DATE: 2002-05-20
PRIOR FILLING DATE: 2002-05-20
PRIOR FILLING DATE: 2002-05-20
PRIOR FILLING DATE: 2002-05-20
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Matches 114;
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                      APPLICANT:
APPLICANT:
APPLICANT:
                                                                                 APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Tchernev, Velizar
APPLICANT: Zhong, Mei
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SOFTWARE: PatentIn version 3.1
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ORGANISM: Homo sapiens
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Anderson, David
Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
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NUMBER OF SEQ ID NOS: 512
SOPTWARE: PatentIn Ver. 2
SEQ ID NO 222
SEQ ID NO 222
LENGTH: 318
TYPE: PRT
ORGANISM: Mus musculus
US-10-055-877-222
                                                                                                                                                                                                                                                                                                    Query Match 29.3%; Score 475; DB 6; Length 318; Best Local Similarity 35.7%; Pred. No. 3.8e-36; Matches 114; Conservative 62; Mismatches 121; Indels
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TITLE OF INVENTION: Novel Polypeptides and Nucleic FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR PILING DATE: 2001-01-23
PRIOR FILING DATE: 2001-01-23
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FILLING DATE: 2001-01-30
APPLICATION NUMBER: 60/272,870
FILING DATE: 2001-03-02
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FILING DATE: 2001-01-25
APPLICATION NUMBER: 60/264,478
FILING DATE: 2001-01-26
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FILING DATE: 2001-01-24
APPLICATION NUMBER: 60/264,117
FILING DATE: 2001-01-25
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                                                                                              RYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKOLPFCRSNILSHSYCLH 184
LAVLKLACADISINVISMGVANVIFLGVPVLFIFVSYIFILSTILRIPSAEGRKKAFSTC
                                                                      RYVAICNPLRYPVVMNKAAYVPMAVSSWVAGGANSLVQISLAVQLPPCGDNVINHFICEI
                                                                                                                                                   FLGNLSFLDICYTTSSVPLVLDGFLTPRKTISFSGCAVOMFLSFAMGATECVLLGMMAFD
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Taupier Jr., Raymond
Pena, Carol
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Shimkets, Richard
Gusev, Vladimir
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Patturajan, Meera
Burgess, Cahterine
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) NOS: 512
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241
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FILE REFERENCE: 21402-251

CURRENT APPLICATION NUMBER: US/10/055,877

CURRENT APPLICATION NUMBER: 60/262,892

PRIOR APPLICATION NUMBER: 60/263,598

PRIOR PILING DATE: 2001-01-19

PRIOR PILING DATE: 2001-01-23

PRIOR PILING DATE: 2001-01-23

PRIOR PILING DATE: 2001-01-23

PRIOR APPLICATION NUMBER: 60/264,117

PRIOR APPLICATION NUMBER: 60/264,117

PRIOR APPLICATION NUMBER: 60/264,117

PRIOR PILING DATE: 2001-01-25

PRIOR PILING DATE: 2001-01-25

PRIOR APPLICATION NUMBER: 60/264,139

PRIOR PILING DATE: 2001-01-25

PRIOR APPLICATION NUMBER: 60/264,478

PRIOR APPLICATION NUMBER: 60/264,351

PRIOR APPLICATION NUMBER: 60/263,351

PRIOR APPLICATION NUMBER: 60/263,351

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-03-02

PRIOR PILING DATE: 2001-03-02

PRIOR APPLICATION NUMBER: 60/272,870

PRIOR APPLICATION NUMBER: 60/272,870

PRIOR APPLICATION NUMBER: 60/272,870

PRIOR APPLICATION NUMBER: 60/275,990

PRIOR APPLICATION NUMBER: 60/275,990
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US-10-055-877-326
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Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 512
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 326
LENGTH: 318
TYPE: PRT
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PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
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APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
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Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
Ratelli, Luca
Kekuda, Ramesh
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Taupier Jr., Raymond
Pena, Carol
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Miller, Charles
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Shimkets, Richard
Gusev, Vladimir
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Burgess, Cahterine
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Mezes, Peter
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Zerhusen, Bryan
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US-10-055-877-326
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Matches 114; Conservative
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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
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APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
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Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
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Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
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Burgess, Cahterine
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Mezes, Peter
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Sequence 12, Application US/11190188
Publication No. US20060035331A1
GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
PRIOR PELICATION NUMBER: 60/272,870
PRIOR FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 28.6%; Score 464; DB 6; Length 319
Best Local Similarity 35.3%; Pred. No. 3.8e-35;
Matches 113; Conservative 64; Mismatches 111; Indels
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LENGTH: 319
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  APPLICANT:
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ORGANISM: Homo sapiens
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APPLICATION NUMBER: 60/264,139
FILING DATE: 2001-01-25
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FILING DATE: 2001-01-24
APPLICATION NUMBER: 60/264,117
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                                                                                           YAO, Monique G.
SHIH, Leo L.
TRIBOULBY, Catheri
LU, Dyung Aina M.
                                                                                                                                                                                           TAL, Preeti
                  KHAN, Farrah A.
POLICKY, Jennifer L.
AU-YOUNG, Janice
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Gaps

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; Sequence 21, Application US/10511538
; Publication No. US20060026700A1
; GENERAL INFORMATION:
APPLICANT: Origene Technologies, Inc
TITLE OF INVENTION: TISSUE SPECIFIC GENES AND GET
FILE REFERENCE: 16U 200 PCT
; CURRENT APPLICATION NUMBER: US/10/511,538
; CURRENT FILING DATE: 2004-10-18
; PRIOR RELIGING DATE: 2002-04-16
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US-10-511-538-21
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PRIOR APPLICATION NUMBER: US 60/411,882
PRIOR FILING DATE: 2002-09-20
PRIOR PELICATION NUMBER: US 60/424,336
PRIOR FILING DATE: 2002-11-07
PRIOR FILING DATE: 2002-11-07
PRIOR APPLICATION NUMBER: US 60/374,823
PRIOR FILING DATE: 2002-04-24
PRIOR FILING DATE: 2002-05-01
PRIOR FILING DATE: 2002-05-01
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PRIOR FILING DATE: 2000-03-03; 2000-03-10; 2000-03-17; 2000-03-20
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PERL Program
SEQ ID NO 12
LENGTH: 319
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APPLICANT:
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyt
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CURRENT APPLICATION NUMBER: US/11/190,188
CURRENT FILING DATE: 2005-07-27
PRIOR APPLICATION NUMBER: US/10/220,382
PRIOR FILING DATE: 2001-03-01
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APPLICANT: BOROWSKY, Mark L.
TITLE OF INVENTION: G-PROTEIN COUPLED RECEPTORS
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Pred. No. 3.8e-35;
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PRIOR APPLICATION NUMBER: US 60/403,648
PRIOR FILING DATE: 2002-08-16
NUMBER OF SEQ ID NOS: 344
SOFTWARE: Patentin version 3.1
SEQ ID NO 21
LENGTH: 319
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Best Local 9
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                               300 EIRORILRLFHVATHA 315
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DMKEAIGRLFHYQKRA
                                                                   FSTCAAHL-AVFLLFFGSVAV-MYLRFSATYSVFWDTAIAVTFVILAPFFNPIIYSLKNK
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Search completed: March Job time : 23 secs 9, 2006, 08:44:24

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Result
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Maximum
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Maximum Match
Listing first
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Perfect score:
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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407.5
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Match
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Gapop 10.0 ,
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2: pir2:*
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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A46750
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olfactory receptor
olfactory receptor
olfactory receptor
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ALIGNMENTS

62

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A;Reference number: A6054
A;Accession: A60547
A;Molecule type: DNA
A;Residues: 1-292 <FEI>A;Cross references: UNIP)
C;Superfamily: olfactory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     odorant receptor 8 - channel catfish
(;Species: Ictalurus punctatus (channel catfish)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change
C;Accession: E45774
R;Ngai, J:; Dowling, M.M.; Buck, L.; Axel, R.; Chess, A.
Cell 72, 657-666, 1993
A;Title: The family of genes encoding odorant receptors in the caption of the caption 
                                                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:Q9PSJ5; UNIPARC:UPI0000042025
A;Experimental source: olfactory epithelium
A;Note: sequence extracted from NCBI backbone (NCBIP:1277-C;Superfamily: olfactory receptor OR14
C;Keywords: olfaction; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Homo sapiens (man)
C;Date: 17-Apr-1993 #sequence revision 17-Apr-1993 #text_change 26-Aug-1999
C;Accession: A60547; A60482
R;Feingold, E.A.; Forget, B.G.
Blood 74, 2118-2186, 1999
A;Title: The breakpoint of a large deletion causing hereditary persistence of fetal hemc A;Reference number: A60547; MUID:90028751; PMID:2478223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Status: preliminary; not compared with conceptual translation A; molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypothetical protein (HPFH breakpoint 3' region) - N;Alternate names: olfactory receptor homolog
                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Residues: 1-313 <NGA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cross-references: UNIPARC:UPI0000178388;Superfamily: olfactory receptor OR14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                            MVDPNGNESSATYFILIGLPGLEBAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEP 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CVFLQFYLLAFFSFFTHRFGSH------IPPYIH 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLN 290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSHV 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CYPLRHANIFTHQLVIQIGTMVVLRAAILVAPCHVLIKCRFQFYHTTVISHSYCEHMAIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ASSIMPKWLGYSGL------MCLKSILIPACFNVVHPHIAGIESGILVAMALDRYVAI 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LVGIPGLESVQCWIGIPFCAIYLIAMIGNSLLLSIIKSERSLHEPLYIFLGMLGATDIAL 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LIGLEGLEEAQFWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIFLCMLSGIDILI
                                    MLAPVQNISFTT-FTLTGFHDLGEWGPILSIPYLLMFLLSSTSNLTLIYLIISQRALHSP 59
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                                                                                                                                                            29.2%; Score 473.5; DB 2; llarity 34.1%; Pred. No. 1.2e-33; Conservative 66; Mismatches 132;
                                                                                                                                                                                                                                                                                                                                                         (NCBIP:127748)
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                                                                                                                                                                                                                       Length 313;
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RESULT 4
A46247
A46247
olfactory receptor OR3 - mouse
olfactory receptor OR3 - mouse
(;Species: Mus musculus (house mouse)
C;Species: 22-5ep-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Date: 22-5ep-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Date: P.; Hermans-Borgmeyer, I.; Artieres-Pin, H.; Beasley, L.; Dionne, V.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Raming, K.; Krieger, J.; Strotmann, J.; Boekhoff, I.; Kubick, Nature 361, 353-356, 1993
A;Title: Cloning and expression of odorant receptors.
A;Reference number: S29707; MUID:93149273; PMID:7678922
A;Accession: S29710
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S29710
S29710
S19710
S19710
S19710
C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Sep-1998
C;Accession: S29710
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A; Residues: 1-307 < RAM>
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            59 FFLASLSLLDALFSTAISPKLIADLLYDQKTISFRACMSQLFIEHLFGGVDIVILVAMAY 118
                                                                                                                                                                                                                                                                                          NSEIKSCMKKLWCKMLHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HODVMKLACDDIRVNVVYGLIVI----ISAIGLDSLLISFSYLLILKTVLGLTREAQAKA 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAFG 241
                                                                                                                                                                                                                                                                                                                                                                                                                     LSTCSSHILVVILFFVPCIFMYARPVYNF-----PIDKCITVFYTITPMLNPLIYTLR 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCL 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAF 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENNNITEFILLGLTQDPDGRKALFVIFF---LIXIVIMMGNLLIVVTVIASPSLGSPMY 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GNESSATYFILIGL---PGLEBAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMY 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMBSTVLLAM 121
                                                                                                                                                                                                                                                                                                                                                         TKEIRORILRLFHVATHA 315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FGTCVSHVCAVFIFYVP--FIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVK 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MSPLIVLACTD---TYPIGLTVIANGGVNCIVIFTLLLG-SYGIILRSLKTQSQEGRRKA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IRQRILRL 308
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLP-VILANIYLLVPPVLNPIVYGVKTKE 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ALDRFFAICRPLYYQKYMGMPNFLKFIIFPVIRNLFFITTIVSWAGKLTFCETNBIDHCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AFDRYVAICHPLRHATVLTLERVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSY 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MCILIGLMAVVDLSMPIFCVPNMLLSFLFNWKGISLVGCLVQMFCIHCAGTFQSTILLWM 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DRYVAICKPLHYLAIMNRRVCITLLIFAWTGGFTHSLIQIVFVYNLPFCGPNVIDHFICD 178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27.9%; Score 453; DB 2; 33.6%; Pred. No. 7.3e-32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     64; Mismatches
H.; Beasley, L.; Dionne, V.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        125; Indels
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             Heinemann,
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Breer,

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R;Buck, L.; Axel, R.

Gell 65, 175-187, 1991

A;Title: A novel multigene family may encode odorant receptors: a mole A;Reference number: A23701; MUID:91191556; PMID:1840504

A;Recession: I23701

A;Retatus: nucleic acid sequence not shown

A;Molecule type: mRNA

A;Residues: 1-312 <BUC>
A;Cross-references: UNIPROT:P23273; UNIPARC:UP10000041F7F; GB:M64391; C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Proc. Natl. Acad. Sci. U.S.A. 89, 8948-8952, 1992
A;Title: Spatial pattern of receptor expression in the olfactory epithelium.
A;Reference number: A46247; MUID:93028384; PMID:1384038
A;Accession: A46247
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: nucleic acid
A;Molecule type: nucleic acid
A;Residues: 1-312 <NEF:
A;Cross-references: UNIPROT:P23275; UNIPARC:UPI0000029735; GB:M84005; NID:g200153; PIDN:
A;Note: sequence extracted from NCBI backbone (NCBIP:115362)
C;Superfamily: olfactory receptor OR14
                                                                                                  원 유
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             olfactory receptor I14 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 09-Jul-2004
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Best Local S
Matches 105
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 125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REAQAKAFGTCVSHVCAVFIFYVPFIGLSMVHRF---SKRRDSPLFVILANIYLLVPFVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KVDNFLCEVPAMIKLACGDTSLNEAVLNGVCTFFTVVPVSVILV--SYCFIAQAVMKIRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ILSHSYCHHQDVMKLACDDIRVN--VVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LLAMAFDRYVAICHFLRHATVLTLPRVTKIGVAAV--VRGAALMAFLFVEIKQLFFCRSN 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LHEPMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEVDSN---SSSGTFILMGVSDHPHLEIIFFAV---ILASYLLTLVGNLTIILLSRLDAR
AICFPLRYTTIMS----TKFCASLVLLLMMLTMTHALLHTL--LIARLSFCEKNVILHFF
                              AICHPLRHATVLTLPRVTKIGVAAVV-----RGAALMAPLPVFIKQLPFCRSNILSHSY 181
                                                                                                          MISGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMBSTVLLAMAFDRYV 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NPLI YTLRNKEVKGALGRLLGKGRGAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NPIVYGVKTKEIRORILRLFHVATHAS 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LVVMAFDRYVAVCRPLHYMTVMN-PRLC-WGLAAISWLGGLGNSVIQSTFTLQLPFCGHR
                                                                         NLSFSDLCFSSVTMPKLLQNMQSQVPSISYTGCLTQLYFFMVFGDMESFLLVVMAYDRYV
                                                                                                                                                 NQTLILBFLLLGLPIPSEYHLLFYALFLAMYLTIILGNLLIIVLVRLDSHLHMPMYLFLS
                                                                                                                                                                                  NESSATYFILIGLPGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIFLC 67
                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                      26.8%;
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34.6%; Pred. No. 1e-30;
                                                                                                                                                                                                                      64;
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                                                                                                                                                                                                                    Score 434.5; DB 2; Length Pred. No. 3e-30; Indels 54; Mismatches 119; Indels
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                                                                                                                                                                                                                                                           Length 312;
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                                                                                                                                                                                                                        25;
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C;Date: 22-Jan-1993 #sequence_revision 22-Jan
C;Accession: E23701
R;Buck, L.; Axel, R.
Cell 65, 175-187, 1991
A;Title: A novel multigene family may encode
A;Reference number: A23701; MUID:91191556; PM
A;Accession: E23701
C;Species: Homo sapiens (man)
C;Date: 14-Oct-1997 #sequence_revision 07-Nov-1997 #text_change
C;Accession: JC5624; 858003
R;Vanderhaeghen, P:; Schurmans, S:; Vassart, G:; Parmentier, M.
Biochem. Biophys. Res. Commun. 237, 283-287, 1997
A;Title: Molecular cloning and chromosomal mapping of olfactory
A;Reference number: JC5624; MUID:97415789; PMID:9268701
A;Accession: JC5624
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B23701
olfactory receptor I3 - rat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  102;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26.6%; Score 431.5; DB 2; 33.2%; Pred. No. 5.4e-30; ative 71; Mismatches 121;
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C;Superfamily: olfactory receptor OK14
C;Keywords: G protein-coupled receptor; transmembrane protein
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                                                                                                                                                                                                                                                                                                      126 YVAICHPLRHATVLTLPRVTKIGVA---AVVRGAALMAPLPVFIKQLPFCRSNILSHSYC 182
                                                                                                                                                                                                                                                                                                                                                                 61 LSNLSFSDLCFSSVTMPKLLQNMRSQDTSIPYGGCLAQTYFFMVFGDMESFLLVAMAYDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLHQDVMKLACDDIRVN-----VVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREA 235
                                                                                                                                                                                                                                                                                                                                                                                                                             LCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDR 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NGTFITQFLLLGLPIPEBHQHLFYALFLV--WYLTTIGNLLIIVLVQLDSQLHTPMYLF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NESSATYFILIGLPGLEBAQ--FWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIF
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TCGSHLSVVSLFYGTIGLYLCP--AGNNSTVKEMVMAMMYTVVTPMLNPFIYSLRNRDM
                                                        TCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKEI 301
                                                                                                                   DLFVLLKLACSDTYINELMIFIMSTLLIIIPFFLIVMSYARIISSILKVPSTQGICKVFS
                                                                                                                                                                            LHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFG 241
                                                                                                                                                                                                                                          YVAICFPLHYTSIMS-PKLCTCLVLLLWMLTTSHAMMHTL--LAARLSFCENNVVLNFFC
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receptor genes expressed

239

185 122 62 65

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C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1994 #sequence_revision 10-Nov.
C;Accession: S29708
R;Raming, K.; Krieger, J.; Strotmann, J.; Boel
Nature 361, 353-356, 1993
A;Title: Cloring and expression of odorant revision of each ratter and the second rate of each r
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A;Residues: 1-319 <VANS
A;Residues: 1-319 <VANS
A;Residues: 1-319 <VANS
A;Cross-references: UNIPROT:Q15622; UNIPARC:UPI0000041DE3; GB:Y10530; NID:g2792017; PID:
R;Vanderhaeghen, P.; Schurmans, S.; Vassart, G.; Parmentier, M.
R;Vanderhaeghen, P.; Schurmans, S.; Vassart, G.; Parmentier, M.
R;Vanderhaeghen, P.; Schurmans, S.; Vassart, G.; Parmentier, M.
A;Description: Male germ cells from several mammalian species express a specific repertor, A;Reference number: S57995
A;Accession: S58003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: mRNA
A;Residues: 126-292 <VAV>
A;Residues: 126-292 <VAV>
A;Crose-references: UNIPARC:UPI000016B10A; EMBL:X89677; NID:g902337; PIDN:CAA61824.1;
A;Crose-references: UNIPARC:UPI000016B10A; EMBL:X89677; NID:g902337; PIDN:CAA61824.1;
C;Genetics:
A;Map position: 19p13.1
C;Function:

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F:58-79/Domain: transmembrane #status predicted <TM2>
F:58-79/Domain: transmembrane #status predicted <TM3>
F:101-120/Domain: transmembrane #status predicted <TM4>
F:140-164/Domain: transmembrane #status predicted <TM4>
F:197-218/Domain: transmembrane #status predicted <TM5>
F:237-260/Domain: transmembrane #status predicted <TM6>
F:272-292/Domain: transmembrane #status predicted <TM7>
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   98; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAF 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GNESSATYFILIGL---PGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ELNQVIQLACSDSFLNHMVIYFTVALLGGGPLTGIL--YSYSKIISSIHAISSAQGKYKA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPV-FIKQLPFCRSNILSHSYC 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DÍKRAL 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FSTCASHLSVVSLFYGAILGVYL--SSAATRNSHSSATASVMYTVVTPMLNPFIYSLRNK 295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FGTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTK 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DRFVAICHPL-HYMVIMNPHLCGLLVLASWTMSALYSLLQILMVVRLSFCTALEIPHFFC
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26.5%; Sc
llarity 31.7%; Pr
Conservative 75;
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Score 429.5; DB 2;
Pred. No. 8.2e-30;
5; Mismatches 119;
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                                                                                                                                                                                                                                                                                                                                                                                                                        Boekhoff, I.;
                                                                                                                                                                                                                                                                                                                      receptors.
PMID:7678922
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       Indels
                                                                     Length 312;
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G45774
G45774
cdorant receptor 202 - channel catfish
C;Species: Ictalurus punctatus (channel catfish)
C;Species: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ringai, J.; Dowling, M.M.; Buck, L.; Axel, R.; Chess, A. Cell 72, 657-666, 1993
A;Title: The family of genes encoding odorant receptors A;Reference number: A45774, MUID:93201590; PMID:7916654
A;Accession: G45774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Experimental source: olfactory epithelium A;Note: sequence extracted from NCBI backbone C;Superfamily: olfactory receptor OR14 C;Keywords: olfaction; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Residues: 1-328 <NGA>
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       문
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           64 IFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAF 123
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Similarity 29.4%; Pred. No. 1.2e-29;
92; Conservative 76; Mismatches 127
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                                                                     GVKTKEIRQRILR
                                                                                                                                              LSTCSSQLIIITLFFLPRCLNYLSSSLGIH-----INADIQILVIMLYSLLPPMINPVIY
                                                                                                                                                                                                                                                                                                                                                                                                                            DRYVAVCNPLRYVNIVKESTILGLCVVSWLLAEPTVLTTVIRATSLPYCASNTVIQCYCD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LSNLSFSDLCFSSVTMFKLLQNMQNQDTSITYTGCLTQMYFSMVFAGMBIFLLVSMAYDR
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       CLRTKEAKECLKR
                                                                                                                                                                                                               FGTCVSHVCAVFIFYVP----FIGLSM-VHRFSKRRDSPLPVILANIYLLVPPVLNPIVY
                                                                                                                                                                                                                                                                                     HVSVTKLACID---RTPYAFPALVSALVMLLTPLAFILFSYGSIIVTVFRTSSTRGRLKT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FGTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTK 299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    YVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQ
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olfactory receptor I15 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 09-Jul-2004
C;Accession: AJ7286
R;Buck, L; Axel, R.
Cell 65, 175-187, 1991
A;Title: A novel multigene family may encode odorant receptors: a molecular basis for ca, Reference number: AZ3701; MUID:91191556; PMID:1840504
A;Accession: AJ7286
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A;Cosss-references: UNIPARC: UNIPARC: UPI0000040A93; GB:M64392; NID:g205845; PIDA
C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
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A; Residues: 1-314 < PAR>
                                                                                                           Matches
                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                             Cross-references: UNIPROT:P23274; UNIPARC:UPI0000040A93; GB:M64392; NID:g205845; PIDN
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NESSATYFILIGLPGLEEAQ--FWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMBSTVLLAMAFDRYV 127
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                                                                                                 26.2%; Score 425.5; DB 2; 33.2%; Pred. No. 1.8e-29; tive 70; Mismatches 114;
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olfactory receptor OR5 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1994 #sequence revision 10-Nov-1995 #text_change 31-Dec-2004
C;Accession: S29707; B37286
C;Accession: S29707; B37286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Buck, L.; Axel, R.
Cell 65, 175-187, 1991
A;Title: A novel multigene family may encode odorant receptors: a molecular basis for A;Reference number: A23701; MUID:91191556; PMID:1840504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Raming, K.; Krieger, J.; Strotmann, J.; Boekhoff, I.; Kubick, Nature 361, 353-356, 1993
A;Title: Cloning and expression of odorant receptors.
A;Reference number: S29707; MUID:93149273; PMID:7678922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPARC:UPI0000E84D3; GB:M64375; NID:g205811; PIDN:AAA41738.1; PID:gC;Keywords: G protein-coupled receptor; membrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: mRNA
A; Residues: 193-236 <BUC>
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A; Residues: 1-314 < RAM>
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301 IR 302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 26.1%; Score 423.5; DB 2; Length 314; Similarity 34.4%; Pred. No. 2.7e-29;
                                                                                                                                                                                                                              CLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAF
                                                                                                                                                                                                                                                                                                       YVAICEPLHYMSIMS----PKLCVSLVVLSWVLTTFHAMLHTLLMARLSFCEDNVIPHFF
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                                                         SSCGSHLSVVSLFYGTVIPLYLCP--SSNNSTVKETVMSLMYTLVTPMLNPFIYSLRNRD
                                                                                                                  GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKE
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A; Molecule type: mRNA
A; Residues: 1-321 < NGA>
A; Cross-references: UNIPROT:Q9PSJ2; UNIPARC:UPI000042021
A; Experimental source: olfactory epithelium
A; Note: sequence extracted from NCBI backbone (NCBIP:12774
C; Superfamily: olfactory receptor OR14
C; Keywords: olfaction; transmembrane protein
                                                                                                                                                                                          olfactory factor OR37 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Sep-1998
C;Accession: S29711
R;Raming, K.; Krieger, J.; Strotmann, J.; Boekhoff, I.; Kubick, S.; Baumstar
Rature 361, 353-356, 1993
A;Title: Cloning and expression of odorant receptors.
A;Reference number: S29707; MUID:93149273; PMID:7678922
A;Accession: S29711
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C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
C;Accession: H45774
R;Ngai, J.; Dowling, M.M.; Buck, L.; Axel, R.; Chess, A.
Cell 72, 657-666, 1993
A;Title: The family of genes encoding odorant receptors in the channel catfish.
A;Reference number: A45774, MUID:93201590; PMID:7916654
A;Accession: H45774
                                                                                                         A;Molecule type: mRNA
A;Residues: 1-305 <RAM>
A;Cross-references: UNIDARC:UPI000003FF7B
C;Superfamily: Olfactory receptor OR14
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S29711
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C;Species: Ictalurus punctatus (channel catfish)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
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                        Conservative
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                                          25.8%;
32.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     320
                        62;
                      Score 418.5; DB 2
Pred. No. 7.2e-29;
52; Mismatches 115
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                                                              DB 2;
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                                                                 Length
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                        31;
                      Gaps
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C(species: Rattus norvegicus (Norway rat)
C(species: Rattus norvegicus (Norway rat)
C(spate: 10-Sep-1999 #sequence revision 10-Sep-1999 #text_change 09-Jul-2004
C(Accession: S51356; S47014
R(stat, U.; Netrasova, E.; Lancet, D.; Natochin, M.
Bur. J. Biochem. 225, 1157-1168, 1994
Bur. J. Biochem. 225, 1157-1168, 1994
A;Title: Olfactory receptor proteins. Expression, characterization and partial purificati
A;Reference number: S51356; MUID:95045546; PMID:7957207

A. Accession. S61356
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A;Residues: 1-309 <GAT:
A;Crose-references: UNIPROT:Q63395; UNIPARC:UPI00000405D8; EMBL:X80671; NID:g517365;
R;Gat, U.; Nekrasova, E.; Lancet, D.; Natochin, M.
submitted to the EMBL Data Library, July 1994
A;Description: Olfactory receptor proteins: expression, characterization and partial
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A;Accession: S47014
A;Status: preliminary
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A; Status: preliminary
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A; Residues: 1-309 < GA2>
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                                                                                                                                                                                                                                       65 SSLSFVDLCYSTVITPKWLVNFLGKKNFITYSECMAQFFFFAIFVVTEGYLLTVMAYDRY
                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
SHSYCLHQDVMKLACDDIRVNVVVGLIVIISAIG--LDSLLISFSYLLILKTVLGL-TRE
                                                                                                                                                                                              VAICHPLRHATVLTLPRVTKIGV-----AAVVRGAALMAPLPVFIKQLPFCRSNIL 177
                                                                                                                                                                                                                                                                                      CMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRY
                                                                                                                                                                                                                                                                                                                                                                  GNESSATYFILIGLPGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIFL
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                                                                                                                                                      VAICRPLLYNVIMS-SRICSLLVLVAFSLGLFSÄVVHTSÄMM-----NLSFCKSYII
                                                                                                                                                                                                                                                                                                                              GNHSAVVVFVLVGLTKQPELLLPLFFLFLVIYVLTVVGNLGMILLIIVSPLLHTPMYYFL
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                                                                   SHYFCDALPLIKLACSNTHLNEL--LIFTIGGLNTLVPTLAVAISYVPIFCSILRIRSSE
                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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A;Gene: GDB:OLFR1
C;Gene: GDB:
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A;Title: The OLFR1 gene encoding the HGMP07E putative olfactory receptor maps to the A;Reference number: A48413; MUID:93251832; PMID:8097991

A;Accession: A48413

A;Mclecule type: DNA

A;Residues: 1-312 < SCH>

A;Coss-references: UNIPROT:P34982; UNIPARC:UPI0000041806; EMBL:X65857; NID:9425220; IA;Note: sequence extracted from NCBI backbone (NCBIN:131483, NCBIP:131485)

C;Genetics:
                                                                                                    RESULT 17
S58066
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C;Species: Homo sapiens (man)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
probable olfactory receptor tpcr33 - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1996 #sequence_revision 19-Apr-1996 #text_change 09-Jul-2004
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R;Schurmans, S.; Muscatelli, F.; Miot, F.; Mattei, M.G.; Vassart, G.; Parmentier, Cytogenet. Cell Genet. 63, 200-204, 1993
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A;Cross-references: UNIPROT:Q9PSJ1; UNIPARC:UPI00000FC739
A;Experimental source: Olfactory epithelium
A;Note: sequence extracted from NCBI backbone (NCBIP:1277
C;Superfamily: Olfactory receptor OR14
C;Keywords: Olfaction; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R.Ngai, J.; Dowling, M.M.; Buck, L.; Axel, R.; Chess, A. Cell 72, 657-666, 1993
A;Title: The family of genes encoding odorant receptors in A;Reference number: A45774; MUID:93201590; PMID:7916634
A;Accession: 145774
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A;Residues: 1-160 <VAN>
A;Cross-references: UNIPROT:Q62336; UNIPARC:UPI0000042009; EMBL:X89685; NID:g902668; PID:
C;Superfamily: olfactory receptor OR14
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R;Vanderhaeghen, P.; Schurmans, S.; Vassart, G.; Parmentier, M.
submitted to the EMBL Data Library, July 1995
A;Description: Male germ cells from several mammalian species express a specific repertoi A;Reference number: S57995
A;Accession: S58066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Ictalurus punctatus (channel catfish)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
C;Accession: 145774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Residues: 1-344 <NGA>
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                                                                                                                   181 YCLHQDVMKLACDDIRVN--VVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQ-A
                                                                                                                                                                                    125 MAYDRVVAICFPLRYNVIVTKRAMTLIIVITWIPSISIIALLVALITRLSFCRSVIINSY 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           246 HYCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVP 286
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238 KAFGTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVK 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  188 MKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLT--REAQAKAFGTCVS 245
                                                                                                                                                                                                                                                                                                                 65 AKYITVFNLALSDLGGSSALIPKLIDTFLFENQVISYEACLANMPFVLFFMTVQSLTLLV 124
                                                                                                                                                                                                                                                                                                                                                                             61 PMYIFLCMLSGIDILISTSSMPXMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 ISNPLRYASVLTNNVIIRIGVAITTRATLSLLPLPFLLKRLNYCPGKILLSHSFCFHADV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h 25.3%; Score 411; DB 2;
Similarity 31.7%; Pred. No. 3.7e-28;
99; Conservative 72; Mismatches 129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NGNESSATY-----FILIGLEGLEEAQFWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHE 60
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                                                                                                                                                                                                                                                      MAPDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVPIKQLPFCRSNILSHS
                                                                 FCDHGPILILAGNDKFINRVMAIGCFVVLDCVPF--LLIIVSYICIGIALMNISHGLERR
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; Pred. No. 1.6e-28;
29; Mismatches 46
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olfactory receptor - dog
C;Species: Canis lupus familiaris (dog)
C;Date: 22-Nov-1993 #sequence_revision 01-Dec-1995 #text_change 09-Jul-2004
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A;Title: Chemoreceptors expressed in taste, A;Reference number: JC5200; MUID:97080538; A;Accession: JC5200
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C;Species: Rattus norvegic
C;Date: 27-Feb-1997 #seque
C;Accession: JC5200; PC430
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;Superfamily: olfactory receptor OR14
;Keywords: olfactiory taste bud; transmembrane protein
;Keywords: olfaction; taste bud; transmembrane predicted <TM1>
;24-47/Domain: transmembrane #status predicted <TM2>
;56-77/Domain: transmembrane #status predicted <TM3>
;199-118/Domain: transmembrane #status predicted <TM4>
;138-162/Domain: transmembrane #status predicted <TM4>
;138-162/Domain: transmembrane #status predicted <TM5>
;136-258/Domain: transmembrane #status predicted <TM6>
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A;Accession: PC4302
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A;Residues: 1-311 <THO1>
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;Residues: 146-153;265-272 <THO2>
;Cross-references: UNIPARC:UPI0000178381; UNIPARC:UPI0000178382
;Experimental source: taste bud
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Date: 27-Feb-1997 #sequence_revision 27-Feb-1997 #text_change 09-Jul-2004
Accession: JC5200; PC4302
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Best Local
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Local Similarity 32.5*; Pred. No. 6.6e-28;
nes 100; Conservative 59; Mismatches 134; Indels 15
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A;Cross-references: UNIPROT:P30955; UNIPARC:UPI0000042014; EMBL:X64996; NID:g890; PIDN:CIC;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
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A;Molecule type: DNA
A;Residues: 1-313 <PAR>
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A;Title: Expression of members of the putative olfactory receptor gene family in mammalian, Reference number: S20571; MUID:92131132; PMID:1370859
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R; Parmentier, M.; L
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                                                                                                                                                                                                                                                                                         VAICHPLRHATVLTLPRVT-----KIGVAAVVRGAALMAPLPVFIKQLPFCRSNILS 178
  IYSLRNKOMKGALRRV
                                         VYGVKTKEIRORILRL
                                                                                                                                                                                                                                                  VAICFPLHYTTIMS-PKLCFSLLVLSWVLTMFHAVLHTLLMA-----RLCFC-ANTIP
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                                                                                  AIGICKVESTCGSHLSVVSLEYGTVIGLYLCP--SANNSTVKETIMAMMYTVVTPMLNPF
                                                                                                                    REAQAKAFGTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPI
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Search completed: March 9, 2006, 08:29:59 Job time : 42 secs

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       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1 MMYDPNGNESSATYFILIGL......KEIRQRILRLFHVATHASEP 318
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1: uniprot_sprot:*
2: uniprot_trembl:*
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(without alignments)
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Q584P5 HUMAN
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Q584P5 HUMAN
Q8V277 MOUSE
Q8WD55 PAT
Q6SHEZ 9PRIM
Q8UBY9 MOUSE
Q8UF94 HUMAN
Q6IF94 HUMAN
Q6IF95 HUMAN
Q6IF95 MOUSE
Q8VH05 MOUSE
Q8VH11 MOUSE
Q8VH17 MOUSE
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Q8VH27 MOUSE
Q8VH27 MOUSE
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Biocceleration
   Q6if93
Q8tif93
Q8tif93
Q584p5
Q584p5
Q66x57
Q6wd57
Q6wd57
Q6wf69
Q9vb59
Q9vb59
Q9vf09
Q9vf09
Q9vf10
Q8vf101
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gorilla gor
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Q7TR86_MOUSE
Q7TR86_MOUSE
Q8VR16_MOUSE
Q8VR16_MOUSE
Q8VR16_MOUSE
Q8VR16_MOUSE
Q8VR116_MOUSE
Q8VR12_MOUSE
Q8VR12_MOUSE
Q8VR11_MOUSE
                          ALIGNMENTS
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Q7trs6 mus musculu
Q8th57 homo sapien
Q8v9x6 mus musculu
Q8v407 mus musculu
Q8v407 mus musculu
Q8v9x8 mus musculu
Q8v9x6 mus musculu
Q8v428 mus musculu
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Q7trq2
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Q6IF93_HUMAN
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W PADMEDIALDS SEVERNES.

W PADMEDIA 1983052; DOI:=10.1073/pnas.0307882100;

W Malnic B., Godfrey P.A., Buck L.B.;

"The human olfactory receptor gene family.";

RI Proc. Natl Acad. Sci. U.S.A. 101:2584-2589(2004).

C -:- MISCELLANEOUS: The sequence shown here is derived from an EMBL/GenBank/DDBJ third party annotation (TPA) entry.

EMBL; BK004369; DAA0476771; -; Genomic_DNA.

DR GO; GO:0004984; F:olfactory receptor activity; IEA.

DR GO; GO:0004984; F:olfactory receptor activity; IEA.

DR GO; GO:0004984; F:olfactory receptor protein signalin. ..; IEA.

GO; GO:0004984; F:olfactory receptor protein signalin. ..; IEA.

DR GO; GO:000186; P:Freceptor activity; IEA.

DR GO; GO:0007165; P:signal transduction; IEA.

DR GO; GO:0007165; P:signal transduction; IEA.

DR InterPro; IPR000725; Olfact receptor.

DR InterPro; IPR000725; Olfact receptor.

DR PANTHER; PTHR11398; Olfact_receptor; 1.

DR PANTHER; PR00237; GFROTEIN_RECEP_F1_1; UNKNOWN_1.

DR PROSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.

KW Receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local !
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                                      OS1E1_HUMAN
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05-JUL-2004 (TREMBLrel. 27, Last sequence update)
05-JUL-2004 (TREMBLrel. 27, Last sequence update)
05-JUL-2004 (TREMBLrel. 27, Last annotation update)
Olfactory receptor OR11-15.
Homo sapiens (Human).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
  10-OCT-2003 (Rel. 42, Created)
                      QSTCB6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q6IF93_HUMAN
Q6IF93;
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317; Conserv
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                                                                                                                                                                                                                GTCVSHVCAVF1FYVPF1GLSMVHRFSKRRDSPLPV1LAN1YLLVPPVLNP1VYGVKTKE
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                                          STANDARD;
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Pred. No. 1.1e-113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94121253465E4B65 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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    R HGNC; HGNC; 1519; OKSIEI.

R InterPro; IPR000725; Olfact_receptor.

R InterPro; IPR000725; Olfact_receptor; 1.

R Pfam; PF00001; 7tm 1; 1.

R Pfam; PF00001; 7tm 1; 1.

R PFNNTS; PR00237; GPCRRHODOPSN.

R PRINTS; PR00245; OLFACTORYR.

R PROSITE; PS00245; OLFACTORYR.

R PROSITE; PS00262; G_PROTEIN_RECEP_F1_2; 1.

R PROSITE; PS00262; G_PROTEIN_RECEP_F1_2; 1.

R PROSITE; PS00262; G_PROTEIN_RECEP_F1_2; 1.

R G_PROTEIN_CEPT_C; 1.

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Bukaryota; Metazoa; C
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AAH22401.1; -; mRNA.
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  Cytoplasmic (Po 2 (Potential). Extracellular (3 (Potential).
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                                                                                                                                                                                                                                                                                                                                sapiens.
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(Potential)

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RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RRA Klausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Korley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickeon M.C.,
RA Blakesley R.W., Touchman J.W., Schmutz J., Myers R.M.,
RGheration and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences ";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).

1- SUBCELLULAR LOCATION: Integral membrane protein.
1- SUBCELLULAR LOCATION: Integral membrane protein.
1- DATABASE: NAME-Human Olfactory Receptor Data Exploratorium
CC. (-1- SUBJERILULAR LOCATION: One of the G-protein coupled receptor 1 family.
1- CC. (-1- DATABASE: NAME-Human Olfactory Receptor Data Exploratorium
                                             This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
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Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases
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13-SEP-2005 (Rel. 48, Last annotation updat
Olfactory receptor 51E1.
as long as its content is in no way modified and this
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Best Local &
Matches 316
weng J., Wang J., Cai Y., Ittmann M., Liu M.;

T "POGR, a prostate-overexpressed G protein-coupled receptor is sensitive biomarker for human prostate cancers.";

submitted (OCT-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; AY75731; AAV54110.1; -; mRNA.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0004872; F:receptor activity; IEA.

GO; GO:0004872; F:receptor activity; IEA.

GO; GO:0007186; P:G-protein coupled receptor protein signalin.

R GO; GO:0007186; P:G-protein coupled receptor protein signalin.

R InterPro; IPR000726; GPCR Rhodpsn.

R InterPro; IPR000725; Olfact_receptor.

R PAINTS; PR00237; GFCRRHODDPSN.

R PRINTS; PR00237; GFCRRHODDPSN.
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QSS45 HUMAN PRELIMINAKY;
QS455;
Q1-FEB-2005 (TrEMBLrel. 29, Created)
O1-FEB-2005 (TREMBLrel. 29, Last sequence update)
O1-FEB-2005 (TREMBLrel. 29, Last annotation update)
O1-FEB-2005 (TrEMBLrel. 29, Last annotation update)
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Mammalia; Eutheria;
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Cytoplasmic (Potential).
N-linked (GlcNAc...) (Potential).
N-linked (GlcNAc...) (Potential).
By similarity.
BC7293AA7FBCA95C CRC64;
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Pred. No. 2.6e-113;
1; Mismatches 0;
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6 (Potential).
Extracellular (Potential)
7 (Potential).
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Best Local Sim
Matches 316;
              TID-GPCE: a novel putative G protein-coupled receptor over prostate cancer and prostate.";

Biochem. Biophys. Res. Commun. 322:239-249(2004).

Biochem. Biophys. Res. Commun. 322:239-249(2004).

Biochem. Biophys. Res. Commun. 322:239-249(2004).

REMBL; AXV998056; AAU07996.1; --; mRNA.

REMBL; MX998056; AAU07996.1; --; mRNA.

REMBL; MX998056; AAU07996.1; --; mRNA.

REMBL; MX998056; F. SIGNATION SECRETORY.

REMBL; MX998057; F. SIGNATION SECRETORY.

REMINITIS, PR000175; GPCRRHODOPSN.

REMINITIS, PR00017; GLECTECEPTORY.

REMINITIS, PR00017; GLECTHOLOPSN.

REMINITIS, PR00017; GPCRRHODOPSN.

REMINITIS, PR00017; GPCRRHOTEIN_RECEP_F1_1; 1.

RECEPTORY.

RECEPTORY.
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PubMed=1531319; DOI=10.1016/j.bbrc.2004.07.106;
Weigle B., Fuessel S., Ebner R., Temme A., Schmitz M.,
Kiessling A., Rieger M.A., Meye A., Bachmann M., Wirth
Rieber E.P.;
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PROSITE; PS50262; G_PROTBIN_RECEP_F1_2; 1.
Receptor.
SEQUENCE 317 AA; 35271 MW; 8C7293AA7FBCA95C CR
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Mammalia; Eutheria;
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01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
01-MAR-2006 (TrEMBLrel. 20, Last annotation update)
01-MAR-2006 (TrEMBLrel. 20, Created)
01-MAR-2006 (TrEMBLrel. 20, Created)
01-MAR-2006 (TrEMBLrel. 20, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-MAR-2003 (TrEMBLrel. 20, Last sequence update)
01-MAR-2005 (TrEMBLrel. 20, Last sequence update)
                     MEDLINE=22974002; PubMed=14611657; DOI=10.1186/gb-2003-4-11-r71; Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A., Walker M., Williams E.M., Trask B.J. tags demonstrate olfactory "Odorant receptor expressed sequence tags demonstrate olfactory expression of over 400 genes, extensive alternate splicing and un expression levels.";
                                                                                                                                                                                                                                                                                                       Young J.M., Priedman C., WILLIAMS Trask B.J.;
"Different evolutionary processes".
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NUCLEOTIDE SEQUENCE.
MEDLINE=21864068; PubMed=11875048; DOI=10.1093/hmg/11.5.535;
Young J.M., Friedman C., Williams E.M., Ross J.A., Tonnes-Priddy L.,
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MEDLINS=21678863; PubMed=11802173;
Zhang X., Firestein S.;
The olfactory receptor gene superfamily
Nat. Neurosci. 5:124-133(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=Ullis (Mouse).
Mus musculus (Mouse).
Metazoa; Chordata;
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NUCLEOTIDE SEQUENCE.
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
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CQSM065;
CQSM065;
COLFEB-2005 (TrEMBLrel. 29, Created)
COLFEB-2005 (TrEMBLrel. 29, Last sequence update)
COLFEB-2005 (TrEMBLrel. 29, Last sequence update)
POLFEB-2005 (TrEMBLrel. 29, Last sequence update)
Prostate overexpressed G protein coupled receptor.
Rattus norvegicus (Rat).
ELMCATYOTA; Metazoa; Chordata; Craniata; Vertebrata; Eutzaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutzaryota; Muridae; Muridae; Rattus.
Muridae; Muridae; Rattus.
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NUCLECTIDE SEQUENCE.
Sanders K.;
Sanders K.;
Compared (JUN-2003)
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SEQUENCE
                                                                                                                                                                                                                                      RAT
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                                                                                                                                                                                                                                                                                                                          RORILRLFLVTTHTSD
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GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0004984; F:olfactory receptor activity; IE3
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0007186; P:receptor activity; IEA.
GO; GO:0007185; P:signal transduction; IEA.
InterPro; IPR000276; GPCR Rhodpsn.
InterPro; IPR000276; GPCR Rhodpsn.
InterPro; IPR000725; Olfact_receptor; 1.
PANTHER; PTHR11398; Olfact_receptor; 1.
PANTHER; PR00237; GPCRRHODOPSN.
PRINTS; PR00237; GPCRRHODOPSN.
PRINTS; PR00237; GPCRHODOPSN.
PROSITE; PS00237; GPCRHODOPSN.
                                                                                                                TCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLFVILANIYLLVPFVLNPIVYGVKTKEI
                                                                         CLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAFG
                                                                                                                                                                          MYIFLCMLSGLDILISTSSMPKMMAIFWENSTTIQFDACLVQMFAIHSLSGMBSTVLLAM
                                                                                                                                                                                        MYIFLCMLSGIDILISTSSMÞKWLAIFWENSTTIQFDACLLQIFAIHSLSGMESTVLLAM
                                                           CLHODVMKLACADIRVNIIYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAFG
                                                                                                                                                                                                                                                                                             Score 1512; DB 2; Pred. No. 1.2e-105; 0; Mismatches 11;
                                                                                                                                                                                                                                                                                                                                                       5A0E59E229852534 CRC64;
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Euteleostomi;

Sciurognathi;

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Best Local S
Matches 291
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InterPro; IPR000725; Olfact_receptor.
PANTHER; PTHR11398; Olfact_receptor; 1.
Pfam; PP00001; 7tm_1; 1.
PRINTS; PR00237; GFCRRHODOPSN.
PRINTS; PR00237; GFCRTORYR.
PROSITE; PS00237; G_PROTEIN_RECEP_P1_1; UNPROSITE; PS00237; G_PROTEIN_RECEP_P1_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weng J., Wang J., Ittmann M., Liu M.;

"POGR, an olfactory G-protein coupled receptor, is overexpressed human prostate and prostate cancers.";

Submitted (NOV-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; AY834218; AAV97881.1; -; mRNA.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0004984; F:olfactory receptor activity; IEA.

GO; GO:0004972; F:receptor activity; IEA.

GO; GO:0007186; P:G-protein coupled receptor protein signalin.

GO; GO:0007185; P:signal transduction; IEA.
                                                                                                                                                                                                                                                                                                          OGSNB2 PPRIM PRELIMINARY; PRT; 223 AA. OGSNB2; OGSNB2; OS-JUL-2004 (TrEMBLrel. 27, Created) 05-JUL-2004 (TrEMBLrel. 27, Last sequence update) 05-JUL-2004 (TrEMBLrel. 27, Last annotation update) 01-JUL-2004 (TrEMBLrel. 27, Last sequence up
                                    WUCLEOTIDE SEQUENCE.
Gilad Y., Wiebe V., Przeworski M., Lancet D., Paabo &
"Loss of Olfactory Receptor Genes Coincides with the
Full Trichromatic Vision in Primates.";
PLOS Biol. 2:0120-0125(2004).
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NUCLEOTIDE SEQUENCE.
STRAIN=BN;
                                                                                                                                                                                                                                            NCBI_TaxID=9593;
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   AY454949; AAR19522.1; -; Genomic_DNA
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ilarity 92.4%; Pred. No. 1.4e-104
Conservative 11; Mismatches 13
                                                                                                                                                                                                                                                                                                                 Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
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Best Local (
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NON_TER
NON_TER
SEQUENCE
Adams M.;
Submitted
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(OBVBV9;

O1-MAR-2002 (TrEMBLrel. 20, Created)

O1-MAR-2002 (TrEMBLrel. 20, Last sequence update)

O1-FEB-2005 (TrEMBLrel. 20, Last sequence update)

O1-FEB-2005 (TrEMBLrel. 20, Last annotation update)

O1-FEB-2005 (TrEMBLrel. 20, Last sequence) (Prostate-

Specific G protein-coupled receptor RAIc) (Mus musculus 0 day neonate

specific G protein-coupled receptor RAIc) (Mus musculus 0 day neonate

specific G protein-coupled receptor NOR18-2, full insert sequence) (Mus musculus 16 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:9630060D12 product:OLFACTORY RECEPTOR MOR18-2, full insert sequence)
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GO; GO:0004994; F:Olfactory receptor act.
GO; GO:0004872; F:receptor act.
GO; GO:0004872; F:receptor act.
GO; GO:0007186; P:G-protein coupled receptor protein signalin..
InterPro; IPR000276; GPCR_Rhodpsn.
InterPro; IPR000275; Olfact_receptor.
PANTHER; PTHR11398; Olfact_receptor;
PANTHER; PRHR11398; Olfact_receptor;
PANTHES; PR00245; OLFACTORYR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Ch
Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
MEDLINE=21676863; PubMed=11802173;
                                                                                                                                                     receptor gene families.";
Hum. Mol. Genet. 11:535-546(2002)
                                                                                                                                                                                                                                                                          Young J.M.,
Trask B.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The olfactory receptor
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                                                                               NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                    "Different evolutionary processes shaped
                                                                                                                                                                                                                                                                                                                                                             MEDLINE=21864068;
                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
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MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
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DR GJ, GO:0016021; C:integral to membrane; TAS.
DR FINTER; DR10727. GFORE C.
DR DR PANTHER; DTHR11398; Olfact receptor; 1.
DR PRINTS, DR10727. GFORENODOEN
                          Query Match
Best Local Similarity
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MEDLINE=20530913; PubMed=11076861; DOI=10:101/gr.152600;
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sequencing pipeline with 384 multicapillary sequencer.";
Genome Res. 10:1757-1771(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN-C57BL/6J; TISSUB=Cerebellum, and Skin; MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100; Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.; Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.; Pubmatarion of Cap-trapper-selected cDNAs prepare full-length cDNA libraries for rapid discovery of new Genome Res. 10:1617-1630(2000).
                                                                        SEQUENCE
                                                                                                                     PRINTS; PR00237; GFORRHODOPSN.
PRINTS; PR00245; OLFACTORYR.
PROSITE; PS00237; G PROTEIN RECEP_F1_1;
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE.
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                                                                              320
                                                                           ξ,
                                                                           35578 MW;
  60.7%;
59.8%;
  Score 985;
Pred. No.
                                                                           7161ACA8F4328959 CRC64;
       5e-66;
                                                                                                                                                    UNKNOWN_1
                          DB 2;
                               Length 320;
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new genes.";
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OSIB2 HUMAN STANDARD,
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
13-SEP-2005 (Rel. 48, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLECTIDE SEQUENCE.;
Xia C., Ma W., Liu M.,
"Identification of a prostate-specific G-protein coupled receptor
(PSGR) that interacts with G alpha subunit in yeast two hybrid
assay.";
                                                                                                                                                                                                                        MEDLINE=21564169; PubMed=11707321; DOI=10.1016/S0378-1119(01)00709-0; Yuan T.T., Toy P., McClary J.A., Lin R.J., Miyamoto N.G., Kretsechmer P.J.; Kretsechmer P.J.; Characterization of an evolutionarily conserved "Cloning and genetic characterization of an evolutionarily conserved human olfactory receptor that is differentially expressed across
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Xu L.L., Stackhouse B.G., Florence K., Zhang W., Shanmugam N., Sesterhenn I.A., Zou Z., Srikantan V., Augustus M., Roschke V., Scarter K., McLeod D.G., Morikantan V., Augustus M., Roschke V., Carter K., McLeod D.G., Morikantan V., Suppet D., Srivastava S.; "PSGR, a novel prostate-specific gene with homology to a G protein-coupled receptor, is overexpressed in prostate cancer."; Cancer Res. 60:6568-6572(2000).
  MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE.
MEDLINE=21002538; PubMed=11118034;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Name=OR51B2; Synonyme=PSGR;
Homo sapiens (Human).
                                                                                                                  NUCLEOTIDE SEQUENCE [LARGE
                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (APR-2001) to
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                                                                                                                  SCALE MRNA].
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Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
AD Distchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
AD Distchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
AD Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
AD Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
AD Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
AD Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
AD Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
AD Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
AD VIIIalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
AD VIIIalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
AD VIIIalon D.K., Wadan A., Young A.C., Shevchenko Y., Bouffard G.G.,
AD AN Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
AD Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
AD Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
AD Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
AN Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
AN Schnerch A., Schein J.B., Jones S.J.M., Marra M.A.,
AN Schnerch A., Schein J.B., Jones S.J.M., Marra M.A.,
AN Generation and initial analysis of more than 15,000 full-length human
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PRINTS; PR00237; GPCRRHODDFSN.

PRINTS; PR00245; OLPACTORYR.

PROSITB; PS00237; G PROTEIN RECEP_F1_1; 1.

PROSITB; PS00237; G PROTEIN RECEP_F1_2; 1.

G-protein coupled receptor; Glycoprotein; Multigene family; Receptor; Sensory transduction; Transducer; Transmembrane.

TOPO_DOM 1 24 Extracellular (Potential).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and mouse cDNA sequences.";

Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).

-!- FUNCTION: Putative odorant receptor.

-!- SUBCELLULAR LOCATION: Integral membrane protein.

-!- TISSUB SPECIFICITY: Exclusively expressed in the prostate.

-!- SIMILARITY: Belongs to the G-protein coupled receptor 1 far.

-!- DATABASE: NAME-Human Olfactory Receptor Data Exploratorium (HORDE);
                                                                                                                                                                                                                                                                                                                                                                                                                                            GO; GO:0016021; C:integral to membrane; NAS.
GO; GO:0004984; F:olfactory receptor activity; NAS.
GO; GO:0007608; P:perception of smell; NAS.
InterPro; IPR000276; GPCR Rhodpsn.
InterPro; IPR000725; Olfact_receptor.
PANTHER; PTHR11398; Olfact_receptor; 1.
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AP369708; AAK38728.1; -; mRNA.
AY333942; AAK37550.1; -; mRNA.
BC020768; AAH20768.1; -; mRNA.
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2 (Potential).
Extracellular (Potential).
3 (Potential).
3 (Potential).
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                                                                   Extracellular (Potential).
5 (Potential).
Cytoplasmic (Potential).
6 (Potential).
Extracellular (Potential).
7 (Potential).
              Cytoplasmic (Potential)
N-linked (GlcNAc. . .)
By similarity.
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Query Match Best Local

Similarity

60.4**%**;

Score Pred.

979; DB 1; No. 1.4e-65;

Matches

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                                                                                                                                                                                                                                                                                                                                                                 Walnice B., Goddrey P.A., Buck L.B.;

If "The human olfactory receptor gene family.";

If Proc. Natl. Acad. Sci. U.S.A. 101:2584-2589 (2004).

If Proc. Natl. Acad. Sci. U.S.A. 101:2584-2589 (2004).

If Roy BK004368; DAA04766.1; -; Genomic_DNA.

If GO; GO:00016021; C:integral to membrane; IEA.

If GO; GO:0004984; F:olfactory receptor activity: IEA.

If GO; GO:0004972; F:receptor activity: IEA.

If GO; GO:0007186; P:G-protein coupled receptor protein signalin. .; IEA.

If GO; GO:0007186; P:G-protein coupled receptor protein signalin. .; IEA.

If GO; GO:0007186; P:G-protein coupled receptor protein signalin. .; IEA.

InterPro; IPR000276; GPCR_Rhodpsn.

InterPro; IPR000725; Olfact_receptor.

IF FAm; PF00001; Ttm 1; 1.

IF FAm; PF00001; Ttm 1; 1.

IF PRINTS; PR00237; GFCRRHODDPSN.

IF PRINTS; PR00237; GFCRRHODDPSN.

IF PRONSITE; PS00237; G_PROTEIN_RECEP_F1_1; UNKNOWN_1.

IF PROSSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.
                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 182
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
01factory receptor OR11-16.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
PubMed=14983052; DOI=10.1073/pnas.0307882100;
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Q6IF94;
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HUMAN
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                                      MLAAIDLALSTSTMPKILALFWFDSREISFEACLTQMPFIHALSAIBSTILLAMAFDRYV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AMPKIS 309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RLFHVA 312
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                                                                                                                                                                                                                                                                                                                                     320 AA;
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                                                                                                                                                                                                                                                                                                                                     35493 MW; 03582CC2AAB6E2C6 CRC64;
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                                                                                                                                                                                                                                   Score 979; DB 2; Length 320; Pred. No. 1.4e-65; Mismatches 67; Indels
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EMBL; AR079864; AAD12761.1; -; mRNA.

R Ensembl; ENSRNOG0000018606; Rattus norvegicus.

R InterPro; IPR000275; GPCR Rhodpsn.

R InterPro; IPR000725; Olfact_receptor.

R PANTHER; PTHR11398; Olfact_receptor; 1.

PEAM; PF00001; 7tm 1; 1.

R PRINTS; PR00237; GPCRRHODOPSN.

R PRINTS; PR00237; GPROTEIN RECEP_F1_2; 1.

R PROSITE; PS00237; G PROTEIN 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            O886.28;
O886.28;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
13-SEP-2005 (Rel. 48, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus (Rat).

Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;

Muroidea; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <del>-</del>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=99131082; PubMed=9932290;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
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MEDINE=21678863; PubMed=11802173;
Zhang X., Firestein S.;
"The olfactory receptor gene superfamily Nat. Neurosci. 5:124-133(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JT 12
9 MOUSE
PRELIMINARY; PRT; 322 AA.
08VP09 MOUSE PRELIMINARY; PRT; 322 AA.
08VP09;
01-MAR-2002 (TrEMBLrel. 20, Created)
01-PAR-2002 (TrEMBLrel. 20, Last sequence update)
01-PER-2005 (TrEMBLrel. 29, Last annotation update)
01-PER-2005 (TrEMBLRel. 29, Cast annotation update)
01-PER-2005 (TrEMBLRel. 29, Last annotation update)
01-PER-2005 (TrEMBLRel. 20, Last annotation update)
                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
MEDLINE-21864068; PubMed-11875048; DOI=10.1093/hmg/l1.5.535;
Young J.M., Friedman C., Williams E.M., Ross J.A., Tonnes-Pr
Trask B.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
MCBI_TaxID=10090;
NUCLEOTIDE SEQUENCE.
MEDLINE=22974002; PubMed=14611657; DOI=10.1186/gb-2003-4-11-r71;
                                                                                                                                                                                           receptor gene fan
Hum. Mol. Genet.
                                                                                                                                                                                                                     "Different evolutionary processes receptor gene families.";
                                                                                         Submitted (JAN-2002)
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6 (Potential).
Extracellular (Potential).
7 (Potential).
7 (Potential).
Cytoplasmic (Potential).
N-linked (GLNAc. . .) (Potential).
                                                                                         EMBL/GenBank/DDBJ
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Pred. No. 2e-65;
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ID OS1D1
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DT 13-SE
DT 13-SE
DT 01fac
GN Name=
OS Homo
OC Eukar
OC Mamma
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REMBL; AY317670; AAP71049.1; -; Genomic_DNA.

REMBL; AY073732; AAL61395.1; -; Genomic_DNA.

REMBL; AY317670; AAL61395.1; -; Genomic_DNA.

RECENTOR:

REMBL; AY317670; AAL701404011.

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Best Local (
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OSIDI HUMAN

STANDARD; PRT; 324 AA.

OBNGF3;

13-SEP-2005 (Rel. 48, Created)

13-SEP-2005 (Rel. 48, Last sequence update)

13-SEP-2005 (Rel. 48, Last sanotation update)

Olfactory receptor 51D1 (Olfactory receptor OR11-14).
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"Odorant receptor expressed sequence tags demonstrate olfactory expression of over 400 genes, extensive alternate splicing and unequal expression levels.";
                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                            Name=OR51D1;
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  NCBI_TaxID=9606;
                                                                                       Homo sapiens (Human)
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                                           Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                           317
                                                                                                                                                                                                                                                                                                                                                                                                  309
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9; Mismatches
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InterPro; IPR000725; Olfact_receptor.
PANTHER; PTHR11398; Olfact_Teceptor; 1.
Pfam; PP00001; 7tm 1; 1.
PRINTS; PR00237; GPCRRHODOPSN.
PRINTS; PR00237; GPCRRHODOPSN.
PRINTS; PR00237; GPROTEIN_RECEP_F1 1; 1.
PROSITE; PS00237; G_PROTEIN_RECEP_F1 2; 1.
G-protein_coupled_receptor; Multigene_family; Olfaction; Receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AB065955; BAC06073.1; -;
EMBL; BK004370; DAA04768.1; -;
Ensembl; ENSG00000197428; Homo
HGNC; HGNC:15193; OR51D1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases
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Suwa M., Sato T., Okouchi I., Arita M., Futami Suwa M., Sato T., Okouchi I., Asai K., Akiyama Y.;
Tsutsumi S., Aburatani H., Asai K., Akiyama Y.;
"Genome-wide discovery and analysis of human se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IDENTIFICATION
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                                             125
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                         RYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLH 184
                                                                                                                  FLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFD 124
                                                                                                                                                                            NGNESSATYFILIGLPGL-EEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYI
                                                                                       FLAMLSTIDLVLSSITMFKMASLFLMGIQEIEFNICLAQMFLIHALSAVESAVLLAMAFD
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1 (Potential).

Cytoplasmic (Potential).

2 (Potential).

Extracellular (Potential).

3 (Potential).

Cytoplasmic (Potential).

Extracellular (Potential).

5 (Potential).

Extracellular (Potential).

5 (Potential).

5 (Potential).

5 (Potential).

6 (Potential).

7 (Potential).

Extracellular (Potential).

8 (Potential).
                                                                                                                                                                                                                                                                 Score 937.5; Pred. No. 1.9e 52; Mismatches
                                                                                                                                                                                                                                                                                                                                                      Cytoplasmic (Potential).
By similarity.
; 52401F88565E3BD1 CRC64;
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Trask B.J.;
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_MOUSE
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                      PF00001;
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EMBL; AY072983; AAL60646.1; -; Genomic DNA.
EMBL; AY317668; AAP71047.1; -; Genomic DNA.
Emsembl; ENSMUSG00000049196; Mus muscullus.
MG1; MG1:3030389; Olffx55.
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0004984; F:olfactory receptor activity; IEA.
GO; GO:0004972; F:receptor activity; IEA.
GO; GO:0007186; P:G-protein coupled receptor protein GO:0007186; P:signal transduction; IEA.
InterPro; IPR000725; Olfact receptor.
PANTHER; PTER11398; Olfact receptor; 1.
prem: DFFMC00725; Olfact receptor; 1.
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01-MAR-2002 (TrEMBLrel. 20, L
01-FEB-2005 (TrEMBLrel. 29, L
01factory receptor MOR10-1 (O
Name=O1fr555;
Mus musculus (Mouse).
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MEDLINE=21676863; PubMed=11802173;
Zhang X., Firestein S.;
The olfactory receptor gene super
                                                                                                                                                                                                                                                                                      Sanders K.; submitted (JUN-2003) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Ch
Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                    expression of over 400 genes, expression levels.";
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Hum. Mol. Genet. 11:535
                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE.
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MEDLINE=22974002; P
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MEDLINE=21864068; Pul
Young J.M., Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Different evolutionary processes shaped the mouse and human olfactory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | SHLCAVLVFYVPLIGLSVVHRLG-GPTSLLHVVMANTYLLLPPVVNPLVYGAKTKBICS
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; 7tm 1; 1.
37; GPCRRHODOPSN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=11875048;
man C., Williams E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PubMed=14611657;
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(Olfactory receptor Olfr555).
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., Tonnes-Priddy L., Ross J.A.,
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B.M., Ross J.A., Tonnes-Pr
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Best Local S
Matches 168
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01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
Olfactory receptor MOR7-2 (Olfactory receptor Olfr577).
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SEQUENCE
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1 MOUSE
QBVH11 MOUSE PRELIMINARY;
                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.

MEDLINB=21676863; Pubmed=11802173;

Zhang X., Firestein S.;

The olfactory receptor gene superfamily
Nat. Neurosci. 5:124-133(2002).
NUCLEOTIDE SEQUENCE.

MEDLINE=22974002; Pubmed=14611657; DOI=10.1186/gb-2003-4-11-r71;
Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A.
Walker M., Williams E.M., Trask B.J.;
"Odorant receptor expressed sequence tags demonstrate olfactory
                                                                                                                                                               NUCLEOTIDE SEQUENCE.
MEDLINE=21864068; PubMed=11875048; DOI=10.1093/hmg/l1.5.535;
YOUNG J.M., Friedman C., Williams E.M., Ross J.A., Tonnes-Priddy
Trask B.J.;
                                                                                                                                                                                                                                                                                                                    Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                            Name=Olfr577;
Mus musculus (Mouse)
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PROSTTE; PS00237; G PROTEIN RECEP F1 1;
PROSITE; PS50262; G PROTEIN RECEP F1 2;
                                                                                                                          receptor gene families.";
Hum. Mol. Genet. 11:535-546(2002)
                                                                          Submitted (JAN-2002) to
                                                                                                   NUCLEOTIDE SEQUENCE
                                                                                                                                                  "Different evolutionary processes
                                                                                                                                                                                                                                                                                                     NCBI_TaxID=10090,
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                                                                          the
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                                                                          EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                            of the mouse.";
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                                                                                                                                                      human olfactory
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RESULT 16
C51G2 HUMA
ID C51G2
AC QBNGK
DT 10-OC
DT 10-SE
DE Olfac
GN Name=
OS Homo
OC Bukar
OC Homo
OC Mamma
OC Homo
OC MOBI
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RP NUCLE
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                                                                                                                                                     O51G2 HUMAN
Q8NGKO;
10-OCT-2003
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GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:001484; F:olfactory receptor activity; IEA.
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0007186; F:G-protein coupled receptor protein signalin.
GO; GO:0007185; P:signal transduction; IEA.
InterPro; IPR000776; GPCR_Rhodpsn.
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PRINTS; PR00237; GECRHODOPSN.

PRINTS; PR00245; OLFACTOXYR.

PROSITE; PS00237; G_PROTEIN_RECEP_F1_1;

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (JUN-2003) to the EMBL/GenBank/DDBJ EMBL; AY072977; AAL60640.1; -; Genomic DNA. EMBL; AY317692; AAP71065.1; -; Genomic DNA. Ensembl; ENSMUSG0000043354; Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression of over 400 genes, extensive alternate splicing and unequal expression levels.";
Genome Biol. 4:R71-R71(2003).
[5]
                                                               Homo sapiens (Human).
Eukaryota; Metazoa; (
Mammalia; Eutheria; E
                                                                                                               10-OCT-2003 (Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last sequ
13-SEP-2005 (Rel. 48, Last anno
Olfactory receptor 51G2.
                                                                                                     Name=OR51G2;
NUCLEOTIDE SEQUENCE.
Suwa M., Sato T., Ok
                                     NCBI_TaxID=9606
                                                                                                                                                                                                HUMAN
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  Okouchi
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                                                                Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
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 Η;
                                                                                                                              sequence update) annotation update)
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 Arita M.,
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I- SUBCELLULAR LOCATION: Integral membrane protein.

I- SIMILARITY: Belongs to the G-protein coupled receptor 1 family.

I- DATABASE: NAME-Human Olfactory Receptor Data Exploratorium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor genes.";
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
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"Genome-wide discovery and analysis of human seven transmembrane helix
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  www="http://bip.weizmann.ac.il/cgi-bin/HORDE/showGene.pl?key=symbol&value=OR510
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                     SHICAVLLFYTPMIGLSVIHRFGKQAPHLVQVVMGFMYLLFPPVMNPIVYSVKTKQIRDR
                                                    SHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKEIRQR
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Cytoplasmic (Potential).
2 (Potential).
Extracellular (Potential).
3 (Potential).
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Pred. No. 4.4e-59;
3; Mismatches 81
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7 (Potential).
7 (Sytoplasmic (Potential).
N-linked (GlCNAc. . .) (Potential).
By similarity.
97DD37E1F38E976C CRC64;
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sapiens.
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RESULT 18
Q8VH17 MOUSE
ID Q8VH17 MOUSE PRELIMINARY;
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I' The human olfactory receptor gene family.";

I' The human olfactory receptor gene family.";

I' Proc. Natl. Acad. Sci. U.S.A. 101:2584-2589(2004).

I' EMBL; BK004285; DAA04683.1; -; Genomic_DNA.

I' GO; GO:0016021; C:integral to membrane; IEA.

I' GO; GO:0004872; F:receptor activity; IEA.

I' GO; GO:0004872; F:receptor activity; IEA.

I' GO; GO:0007186; P:G-protein coupled receptor protein signalin. . .;

I' GO; GO:0007186; P:G-protein coupled receptor protein signalin. . .;

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Q61FH7;
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Mammalia; Eutheria;
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GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0004984; F:olfactory receptor activity; IEA.
GO; GO:0004986; F:olfactory receptor activity; IEA.
GO; GO:000486; P:G-protein coupled receptor protein coupled receptor protein coupled receptor protein go; GO:0007165; P:signal transduction; IEA.

InterPro; IPR000276; GFCR Rhodpsn.

InterPro; IPR0000276; GFCR Rhodpsn.

InterPro; PTHR11398; Olfact receptor; 1.
PFAMN; PR00001; 7tm 1; 1.
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PROSITE; PS00237; GFCRHODDESN.
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Submitted (JUN-2003) to the EMBL/GenBank/DDBJ dat Submitted (JUN-2003) to the EMBL/GenBank/DDBJ dat Submitted (JUN-2003); ALEGO (1: -; Genomic DNA. EMBL; AY317693; AAP71066.1; -; Genomic DNA. Emsembl; ENSWUSG00000045792; Mus musculus. PMGI; MGI; GO:GO:0016021; C:integral to membrane; IEA. GO; GO:0004904; F:olfactory receptor activity; IIA. GO; GO:0004904; F:olfactory receptor activity; IEA. GO; GO:0004872; F:receptor activity; IEA. GO; GO:0007165; P:G-protein coupled receptor GO; GO:0007165; P:signal transduction; IEA.
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MEDLINS=21676863; Pubmed=11802173;
Zhang X., Firestein S.;
"The olfactory receptor gene superfamily Nat. Neurosci. 5:124-133(2002).
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01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
01factory receptor MOR7-1 (Olfactory receptor Olf
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Mammalia; Eutheria;
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Hum. Mol. Genet. 11:535-546(2002).
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Q8VGZE MOUSE PRELIMINARY;
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DT 01-MAR-2002 (TREMBLIFEL 20, LE
DT 01-MAR-2002 (TREMBLIFEL 20, LE
DT 01-FEB-2005 (MOUSE).

OX MUS mUSCULUS (MOUSE).

OX MUS mUSCULUS (MOUSE).

OX MUCHANTIAE; MUTINAE; MUS.

OX NCBI_TAXID=10090;

RN VICLEOTIDE SEQUENCE.
RN MEDLINE=21864068; PubMed=1180;
RN WEDLINE=21864068; PubMed=1187;
RN MEDLINE=21864068; PubMed=1187;
RN MEDLINE=21864068; PubMed=1187;
RN MEDLINE=21864068; PubMed=1187;
RN MEDLINE=21864068; PubMed=1187;
RN MUCLEOTIDE SEQUENCE.
RN MEDLINE=22974002; PubMed=1461;
RN MUCLEOTIDE SEQUENCE.
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RN MUCLEOTIDE SEQUENCE.
RN MEDLINE=23974002; PubMed=1461;
RN MEDLINE=2397400
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     Submitted (UUN-2003) to the EMBL/GenBank/DDBJ database EMBL; AY0772994; AAL60657.1; -; Genomic DNA. EMBL; AY317676; AAP71055.1; -; Genomic DNA. Ensembl; ENSMUSGO00000051711; Mus muscullus. MGI; MGI:3030395; Olfr561.

GO; GO:0016021; C:integral to membrane; IEA. GO; GO:0004984; F:olfactory receptor activity; IEA. GO; GO:0004982; F:receptor activity; IEA. GO; GO:0007186; P:G-protein coupled receptor protein s GO; GO:0007186; P:signal transduction; IEA. InterPro; IPR000725; Olfact receptor. PANTHER; PTHR11398; Olfact receptor; 1.
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MEDLINE=21676863; PubMed=11802173;
Zhang X., Firestein S.;
The olfactory receptor gene super
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
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MEDLINE=21864068; PubMed=11875048;
Young J.M., Friedman C., Williams E
Trask B.J.;
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01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-PEB-2005 (TrEMBLrel. 29, Last annotation update)
01factory receptor MOR30-1 (Olfactory receptor Olfr
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PRINTS; PR00237; GPCRRHODOPSN.

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PROSITE; PS00237; G PROTEIN RECEP_F1_1;

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;

Receptor.
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ZZ MOUSE
QBVGZZ MOUSE PRELIMINARY;
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MEDLINE=21864068, PubMed=11875048, DOI=10.1093/hmg/11.5.535;
Young J.M., Friedman C., Williams E.M., Ross J.A., Tonnes-Priddy L.,
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  MEDLINE=22974002; PubMed=14611657; DOI=10.1186/gb-2003-4-11-r71;
Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A.
                                                                                         Submitted (JAN-2002)
                                                                                                                                                                                                   "Different evolutionary processes receptor gene families.";
                                                                                                                                                                                                                                              Young J.M.,
Trask B.J.;
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GO; GO:0016021; C:integral to membrane; IRA.
GO; GO:0004984; F:olfactory receptor activity; IEA.
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0007186; P:G-protein coupled receptor prot.
GO; GO:0007165; P:signal transduction; IEA.
InterPro; IPR000276; GPCR Rhodgen.
InterPro; IPR000276; GPCR Rhodgen.
InterPro; IPR000725; Olfact receptor.
PANTHER; PTHR11398; Olfact receptor; 1.
Pfam; PF00001; 7tm 1; 1.
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SEQUENCE
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PRINTS; PR00245; OLFACTORYR.

PROSITE; PS00237; G PROTEIN RECEP_F1_1;

PROSITE; PS50262; G PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (JUN-2003) to the EMBL/GenBank/DDBJ EMBL; AY072998; AAL60661.1; -; Genomic_DNA. EMBL; AY317684; AAP71059.1; -; Genomic_DNA. Ensembl; ENSMUSG0000062142; Mus musculus.
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"Odorant receptor expressed sequence tags demonstrate olfactory expression of over 400 genes, extensive alternate splicing and expression levels.";
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US-09-465-901-3
US-08-467-948A-2
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APPLICANT: Wang, Aljun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 21012.427C23
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
TYPE: PRT
TYPE: PRT
RESULT 2
US-10-012-896-920
; Sequence 920, Application US/10012896
; Patent No. 6943236
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Patent No. 6800746
GENERAL INFORMATION:
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Local Similarity 99.7%;
hes 317; Conservative
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Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
                                                                                       IRQRILRLFHVATHASEP 318
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Mitcham, Jennifer L.
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Dillon, Davin
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Pred. No. 1e-139;
1; Mismatches 0;
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RESULT 3
US-09-680-728-2
; Sequence 2, Application U
; Patent No. 6790631
; GENERAL INFORMATION:
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-012-896-920
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C27
CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 1011
SOFTWARE: FaetSEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
THERAPY
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Best Local Similarity 99.7%;
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APPLICANT:
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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                                                                                       GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKE 300
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                   IRQRILRLFHVATHASEP 318
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IRQRILRLFHVATHASEP
                                                                    GTCVSHVCAVF1FYVPF1GLSMVHRFSKRRDSPLPV1LAN1YLLVPPVLNP1VYGVKTKE
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Day, Craig H.
Vedvick, Thomas (
Carter, Darrick
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Wantanabe, Yoshihiro
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Houghton, Raymond L.
Vinals de Bassols, Carlota
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Hepler, William T.
Henderson, Robert A.
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Kalos, Michael D.
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Stolk, John A.
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Pred. No. 1e-139;
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US/09680728

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APPLICANT: Steve Chappell Mitchell
APPLICANT: Douglas C. Saffran
ITILE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR
ITILE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
ITILE OF INVENTION UPMBER: US/9/680,728
CURRENT APPLICATION NUMBER: US/9/680,728
CURRENT PILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR PILING DATE: 1999-10-05
NUMBER OF SEQ ID NOS: 50
SOFTWARE: FastSFO f
SEO ID NOS: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; SEQ ID NO 2
; LENGTH: 317
; TYPE: PRT
; ORGANISM: Homo Sapiens
US-09-680-728-2
FILE REFERENCE: 511582002410
CURRENT APPLICATION NUMBER: US/10/017,066A
CURRENT FILING DATE: 2002-05-28
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR FILING DATE: 1999-10-05
PRIOR FILING DATE: 50
NUMBER OF SEQ ID NOS: 50
SOPTWARE: PastSEQ for Windows Version 4.0
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APPLICANT: Arthur B. Raitano

APPLICANT: Daniel E.H. Afar

APPLICANT: Aya Jakobovite

APPLICANT: Mary Faris

APPLICANT: Rene S. Hubert

APPLICANT: Steve Chappell Mitchell

APPLICANT: Douglas C. Saffran

TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR

TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER.
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/10017066A Patent No. 6838258
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Best Local
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Local Similarity 100.0%; Pred. No. 1.5e-139;
hes 317; Conservative 0; Mismatches 0;
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APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Marlocker, Susan Louise
APPLICANT: Jiang Yuqui
APPLICANT: Kalos, Michael
APPLICANT: Fanger, Gary
APPLICANT: Bolk, John
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
FILE OF INVENTION NUMBER: US/09/439,313
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 527
LENGTH: 320
TYDE: DET
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US-09-439-313-527
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                                                                                                           Query Match 60.4%; Score 979; DB 2; Length 320; Best Local Similarity 59.5%; Pred. No. 2.1e-81; Matches 182; Conservative 55; Mismatches 67; Indels
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TYPE: PRT
ORGANISM: Homo Sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62 MYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMBSTVLLAM
68 MLSGIDILISTSSMPKWLAIFWENSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYV 127
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                                                                          8 NESSATYFILIGLEGLEGAGEWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHEFMYIFLC
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                                     CLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAFG
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US-09-636-215-527
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US-09-636-215-527
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Best Local S
Matches 182
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Dillon, Davin C.
Mitcham, Jennifer L.
MKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSH
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Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
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                                                    AICHPLRHAAVLNUTVTAQIGIVAVVRGSLFFFPLPLLIKRLAFCHSNVLSHSYCVHQDV
                                                                                             AICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQDV 187
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                                                                                                                                                                                                                                                                                                Conservative
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US-09-685-166A-527
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US-09-685-166A-527
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Best Local (
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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                                      307 RLFHVA 312
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Similarity 59.5%; Pred. No. 2.1e-81;
82; Conservative 55; Mismatches 67
                                                                                                   MKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSH
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      AMPKIS
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Henderson, Robert A
Kalos, Michael D.
Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
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US-09-679-426-527

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APPLICANT: Skelky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THI
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C20
CURRENT APPLICATION NUMBER: US/09/679,426
CURRENT FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 895
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 527
LENGTH: 320
                      RESULT 9
US-09-143-527
J Sequence 527, Application US/09759143
Patent No. 6800746
Patent No. 6800746
GENERAL INFORMATION
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Patent No. 6759515
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
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Best Local Similarity 59.5
Matches 182; Conservative
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Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
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Henderson, Robe
Kalos, Michael
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Carter, Darrick
Xu, Jiangchun
Dillon, Davin
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Mitcham, Jennifer L.
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GENERAL INFORMATION:
APPLICANT: XL, Jiang
APPLICANT: Dillon, II
APPLICANT: Mitcham,
APPLICANT: Harlocker
APPLICANT: Harlocker
APPLICANT: Henderson
APPLICANT: Kalos, Mi
APPLICANT: Fanger, G
APPLICANT: Retter, W
APPLICANT: Stolk, Jc
APPLICANT: Day, Crai

Jiang, Yuqui Henderson, Robert Kalos, Michael D.

Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
Day, Craig H.

Patent No. Sequence 527,

27, Application US/09651236 6818751

Xu, Jiangchun Dillon, Davin C.

Mitcham, Jennifer

Harlocker,

Susan L.

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RESULT 10
US-09-651-236-527
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; ORGANISM: Homo sapiens
US-09-759-143-527
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SEQ ID NO 527
LENGTH: 320
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APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C23
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
NUMBER OF SEQ ID NOS: 934
NUMBER OF SEQ ID NOS: 934
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Vedvick,
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APPLICANT: Wang, Aljun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REPERENCE: 210121.42718C18
CURRENT APPLICATION NUMBER: US/09/651,236
CURRENT APPLICATION NUMBER: US/09/651,236
CURRENT FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 865
SOPTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 527
LENGTH: 320
; LENGTH: 320
; TYPE: PRT
; ORGANISM: Homo E
US-09-968-033C-4
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-651-236-527
                                                                                                                       FILE REFERENCE: PF180P1
CURRENT APPLICATION NUMBER: US/09/968,033C
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: 60/237,275
PRIOR APPLICATION NUMBER: 60/237,275
PRIOR APPLICATION NUMBER: 09/339,115
PRIOR FILING DATE: 1999-06-24
PRIOR FILING DATE: 1999-06-24
PRIOR APPLICATION NUMBER: 09/053,303
PRIOR HILING DATE: 1998-04-01
PRIOR APPLICATION NUMBER: 08/465,980
PRIOR APPLICATION NUMBER: 08/465,980
PRIOR FILING DATE: 1995-06-06
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                                                 NUMBER OF SEQ ID NOS: 12
SOPTWARE: PatentIn Ver. :
SEQ ID NO 4
LENGTH: 320
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APPLICANT: Soppet et al.
TITLE OF INVENTION: Human Prostate Specific G-Protein Receptor HPRAJ70
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Best Local Similarity 59.5%; Pred, No. 2.1e-81;
Matches 182; Conservative 55; Mismatches 67;
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Carter, Darrick
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US-09-657-279-527
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APPLICANT: Xu, Jiang
APPLICANT: Dillon, II
APPLICANT: Mitcham,
APPLICANT: Harlocker
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SEQ ID NO 527
LENGTH: 320
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Matches
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APPLICANT:
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APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C19
CURRENT APPLICATION NUMBER: US/09/657,279
CURRENT FILING DATE: 2000-09-06
                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
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Dillon, Davin C.
Mitcham, Jennifer L.
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                                                      60.4%; Score 979; DB 2; Length 320; Ilarity 59.5%; Pred. No. 2.1e-81; Conservative 55; Mismatches 67; Indels
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vick, Thomas (
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                                                                                                                                                      , TYPE: PRT
; ORGANISM: Homo sapiens
US-10-012-896-527
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CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 1011
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 527
LENGTH: 320
                                                                                              Query Match 60.4%; Score 979; DB 2; I Best Local Similarity 59.5%; Pred. No. 2.1e-81; Matches 182; Conservative 55; Mismatches 67;
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APPLICANT:
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
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                                          RLFHVA 312
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McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
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Mitcham, Jennifer L.
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Hepler, William T.
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Kalos, Michael D.
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                                                                                                                          Length 320;
                                                                                                 Indels
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SEQ ID NO 3
LENGTH: 320
TYPE: PRT
ORGANISM: Rat Protein
US-09-680-728-3
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US-09-680-728-3
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Best Local Similarity
Matches 182; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Steve Chappell Mitchell
APPLICANT: Douglas C. Saffran
TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
FILE REFERENCE: 129.24USU1
CURRENT APPLICATION NUMBER: US/09/680,728
CURRENT FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR FILING DATE: 1999-10-05
NUMBER OF SEQ ID NOS: 50
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APPLICANT:
APPLICANT:
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  304 AMFKIS
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                                                                                                                                188 MKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSH 246
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                             307 RLFHVA 312
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Daniel E.H. Afar
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                                                                                                                                                                                                                                                                                                                                       60.2%; Score 977; DB 2; Length 320; llarity 59.5%; Pred. No. 3.3e-81; Conservative 53; Mismatches 69; Indels
    309
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APPLICANT: Arthur B. Raitano
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
APPLICANT: Aya Jakobovits
APPLICANT: Mary Faris
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mit
                 Sequence 2, Application US/09968033C
Patent No. 6824993
GENERAL INFORMATION:
APPLICANT: SODDET et al.
TITLE OF INVENTION: Human Prostate Specific G-Protein Receptor HPRAJ70
FILE REFERENCE: PF180P1
CURRENT APPLICATION NUMBER: US/09/968,033C
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: 60/237,275
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
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SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 3
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CURRENT APPLICATION NUMBER: US/10/017,066A
CURRENT FILING DATE: 2002-05-28
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR PILING DATE: 1999-10-05
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APPLICANT: Steve Chappell Mitchell
APPLICANT: Douglas C. Saffran
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR
TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 320
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APPLICATION NUMBER: 09/339,115
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; NAME/KEY: SITE
; LOCATION: (192)
; OTHER INFORMATION: Xaa equals any amino acid
US-09-968-033C-2
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US-08-465-980-2
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west Local Similarity 59.4%;
Matches 177; Conservation
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Patent No.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve:
CURRENT APPLICATION DATA:
CURRENT APPLICATION UNMBER: US/08/465,980
FILING DATE: 06-JUN-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: PETTATO, GREGOTY D.
REGISTRATION NUMBER: 36,134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1999-06-24
PRIOR APPLICATION NUMBER: 09/053,303
PRIOR FILING DATE: 1998-04-01
PRIOR APPLICATION NUMBER: 08/465,980
PRIOR FILING DATE: 1995-06-06
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver: 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Soppet, Daniel R.
APPLICANT: Li, Yi
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: HUMAN G-PROTEIN RECEPTOR HPRAJ70
NUMBER OF SEQUENCES: 8
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                                                                                                                                                                                                                                                                                               CITY: Roseland
STATE: New Jersey
                                                                                                                                                                                                                                                                         COUNTRY:
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o. 5756309
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Pred. No. 3.7e-80;
                                                                                                                                                                                                                                                                                                                                                                                    BAIN, GILFILLAN, CECCHI,
                                                                                                                                                      Version #1.30
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REFERENCE/DOCKET NUMBER: 325800-446
TELECOMMUNICATION INFORMATION:

201-994-1700

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TELEPHONE:

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APPLICANT: Soppet,
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 320 amino acid
                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
FILING DATE: 06-UUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                     CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: HUMAN G-PROTEIN RECEPTOR HPRAJ70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
NAME: Ferraro, Gregory D. A
REGISTATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 32
IELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                            CITY: Roseland
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEB: CARELLA, BYRNE, BAIN, GILFILLAN, CECCHI, ADDRESSEB: STUART & OLSTEIN
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ilarity 59.4%;
Conservative 5
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Li, Yi
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                                                                                                              08/465,980
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                   325800-446
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Pred. No. 4.5e-80;
                                                                                                                                                                                                                             Version #1.30
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RESULT 19
US-09-339-115-2
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 320 amino acids
TYPE: amino acid
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Patent No. 6372891
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Best Local Similarity
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                                                                                                                             FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
NAME: Ferraro, Gregory D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 325800-446
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 8
                                                                               ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 320 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                              CITY: Roseland
STATE: New Jersey
                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                               APPLICATION NUMBER: US/09/339,115
                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: CARELLA, BYRNE, I
ADDRESSEE: STUART & OLSTEIN
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Li, Yi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Daniel R.
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                                                                                                                                                                                                                                                                                                                                                                                                BAIN, GILFILLAN, CECCHI
                                                                                                                                                                                                                   Version #1.30
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RESULT 20
US-09-680-728-4
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; TYPE: PRT
; ORGANISM: Homo Sapiens
US-09-680-728-4
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Best Local Similarity 59.4%,
Matches 177; Conservative
                                                                                                                                                                                                Query Match 59.5*; Score 964.5; DB 2; Length 320; Best Local Similarity 59.4*; Pred. No. 4.5e-80; Matches 177; Conservative 54; Mismatches 66; Indels 1
                                                                                                                                                                                                                                                                                                                                                        SOFTWARE:
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: René S. Hubert
APPLICANT: Steve Chappell Mitchell
APPLICANT: Steve Chappell Mitchell
APPLICANT: Douglas C. Saffran
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR
TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
FILE REFERENCE: 129 24 USU1
CURRENT APPLICATION NUMBER: US/09/680,728
CURRENT FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR PILLORIUN UNUMBER: 60/157,902
PRIOR PILLORIUN UNUMBER: 50/157,902
PRIOR PILLORIUN UNUMBER: 50/10-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Arthur B. Raitano
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 320 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                136 ATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQDVMKLACDDI 195
  136 ATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQDVMKLACDDI 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              132 AAVLNNTVTAQIGIVAVVRGSLFFFPLPLLIKRLAFCHSNVLSHSYCVHQDVMKLAYADT 191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            76 ISTSSMPKMLAIFWENSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYVAICHPLRH 135
                                        LSTSTMPKILALFWFDSREISIEACLTQMFFIHALSAIESTILLAMAFDRYVAICHPLRH 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VPPIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKBIRQRILRLFHVA 312
                                                                                                                                                                                                                                                                                                                                                                          FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VPLIGLSVVHRFGNSLHPIVRVVMGDIYLLLPPVINPIIYGAKTKQIRTRVLAMFKIS 309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mary Faris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 59.5%; Score 964.5; DB 2;
59.4%; Pred. No. 4.5e-80;
tive 54; Mismatches 66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 320;
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                           255 VPFIGLSMYHRFSKRRDSPLFVILANIYLLVPFVLNFIVYGVKTKEIRQRILRLFHVA 312
                                                                                                                         196 RVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSHVCAVFIFY 254
                                                                                                                                                                         132 AAVLNUTVTAQIGIVAVVRGSLFFFPLPLLIKRLAFCHSNVLSHSYCVHQDVMKLAYADT 191
                                                                                    192 LPNVVÝGLTAILLVMGVDVMFÍSLSÝFLIIRTVLQLPSKSERÁKAFGTCVSHIGVVLAFÝ 251
VPLIGLSVVHRFGNSLHPIVRVVMGDIYLLLPPVINPIIYGAXTKQIRTRVLAMFKIS 309
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Search completed: March 9, 2006, 08:30:51 Job time : 47 secs